

OM protein - protein search, using sw model
 Run on: August 26, 2005, 19:04:47 ; Search time 171 Seconds
 (without alignments)
 726.024 Million cell updates/sec

Title: US-10-767-374-2
 Perfect score: 1688
 Sequence: 1 MGILLGLLLGLHVTYVGR.....AYIMLCRTSQEHVYEAAR 321
 Scoring table: BLOSUM62
 Gapop 10.0 , Gapext 0.5
 Searched: 2105692 seqs, 386760381 residues
 Total number of hits satisfying chosen parameters: 2105692
 Minimum DB seq length: 0
 Maximum DB seq length: 2000000000
 Post-processing: Minimum Match 0%
 Maximum Match 100%

Database : A_Geneseq_18Dec04.*
 Listing first 1500 summaries
 1: Geneseqp1980s.*
 2: Geneseqp1990s.*
 3: Geneseqp2000s.*
 4: Geneseqp2001s.*
 5: Geneseqp2002s.*
 6: Geneseqp2003as.*
 7: Geneseqp2003bs.*
 8: Geneseqp2004s.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

No.	Score	Match	Length	DB	ID	Description
RESULT 1						
ID	AA23322	standard; protein; 321 AA.				
DE	Amino acid sequence of the PRO362 polypeptide.					
PN	W09927098-A2.					
PD	03-JUN-1999.					
PA	(GETH) GENENTECH INC.					
Query Match	100.0%;	Score 1688;	DB 2;	Length 321;		
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;				
RESULT 2						
ID	AA41691	standard; protein; 321 AA.				
DE	Human PRO 362 protein sequence.					
PN	W09946281-A2.					
PD	16-SEP-1999.					
PA	(GETH) GENENTECH INC.					
Query Match	100.0%;	Score 1688;	DB 2;	Length 321;		
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;				
RESULT 3						
ID	AAB33429	standard; protein; 321 AA.				
DE	Human PRO362 protein UNQ317 SEQ ID NO:80.					
PN	W020053758-A2.					
PD	14-SEP-2000.					
PA	(GETH) GENENTECH INC.					
Query Match	100.0%;	Score 1688;	DB 3;	Length 321;		
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;				
RESULT 4						
ID	AAB44247	standard; protein; 321 AA.				
DE	Human PRO362 (UNQ317) protein sequence SEQ ID NO:52.					
PN	W020053756-A2.					
PD	14-SEP-2000.					
PA	(GETH) GENENTECH INC.					
Query Match	100.0%;	Score 1688;	DB 3;	Length 321;		
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;				
RESULT 5						
ID	AA95346	standard; protein; 321 AA.				
DE	Human PRO362 antitumour protein.					
PN	W0200037638-A2.					
PD	29-JUN-2000.					
PA	(GETH) GENENTECH INC.					
Query Match	100.0%;	Score 1688;	DB 3;	Length 321;		
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;				
RESULT 6						
ID	AAB24047	standard; protein; 321 AA.				

DE Human PRO362 protein sequence SEQ ID NO:14.
 PN W020053754-A1.
 PD 14-SEP-2000.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 3; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 7
 ID ABO25193 standard; protein; 321 AA.
 DE Novel human secreted and transmembrane protein PRO362.
 PN US2003050239-A1.
 PD 13-MAR-2003.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 6; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 8
 ID ABU72199 standard; protein; 321 AA.
 DE Novel human secreted and transmembrane protein PRO362.
 PN US2002192706-A1.
 PD 19-DEC-2002.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 6; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 9
 ID AB084879 standard; protein; 321 AA.
 DE Human secreted and transmembrane polypeptide PRO362.
 PN US2002177553-A1.
 PD 28-NOV-2002.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 6; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 10
 ID ABU61077 standard; protein; 321 AA.
 DE Human PRO362 polypeptide.
 PN US2002169284-A1.
 PD 14-NOV-2002.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 6; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 11
 ID ABU80346 standard; protein; 321 AA.
 DE Human secreted/transmembrane protein PRO362.
 PN US2003004102-A1.
 PD 02-JAN-2003.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 6; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 12
 ID ABU07737 standard; protein; 321 AA.
 DE Human A-33 related antigen PRO362.
 PN US2002182206-A1.
 PD 05-DEC-2002.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 6; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 13
 ID ADA24591 standard; protein; 321 AA.
 DE Novel human secreted and transmembrane protein PRO362.
 PN US2003050241-A1.
 PD 13-MAR-2003.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 6; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 14
 ID ABO19648 standard; protein; 321 AA.
 DE Novel human secreted and transmembrane protein PRO362.
 PN US2003050240-A1.
 PD 13-MAR-2003.
 PA (GETH) GENENTECH INC.
 Query Match 100.0%; Score 1688; DB 6; Length 321;
 Best Local Similarity 100.0%; Pred. No. 9.4e-136;

RESULT 15
 ID ADA12252 standard; protein; 321 AA.
 DE Human secreted/transmembrane polypeptide PRO362.

PN US2003055216-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 6; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 16
ID ABO19539 standard; protein; 321 AA.
DE Novel human secreted and transmembrane polypeptide #7.
PN US2003049633-A1.
PD 13-MAR-2003.
Query Match 100.0%; Score 1688; DB 6; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 17
ID ADB73558 standard; protein; 321 AA.
DE Human PRO polypeptide #7.
PN US2003045462-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 18
ID ADB76274 standard; protein; 321 AA.
DE Human PRO polypeptide #7.
PN US2003083248-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 19
ID ADC43700 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003054986-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 20
ID ADC61460 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003049684-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 21
ID ADC63424 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003054405-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 22
ID ADC65624 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003060406-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 23
ID ADC68648 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003064407-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 24
ID ADC62708 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003068648-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.

PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 25
ID ADC67773 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003069178-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 26
ID ADC41093 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003072745-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 27
ID ADC67148 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003073131-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 28
ID ADC62084 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003073624-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 29
ID ADC41717 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003104998-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 30
ID ADC78883 standard; protein; 321 AA.
DE Human PRO protein #56.
PN WO2003034984-A2.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 31
ID ADB49086 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003096744-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 32
ID ADE35140 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003203434-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 7; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 33
ID ADE16254 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003203435-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.

Query Match	100.0%;	Score 1688;	DB 7;	Length 321;
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;		
RESULT 34				
ID	ADH72869 standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.			
PN	US2003203436-A1.			
PD	30-OCT-2003.			
PA	(GETH) GENENTECH INC.			
Query Match	100.0%;	Score 1688;	DB 7;	Length 321;
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;		
RESULT 35				
ID	ADD72227 standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.			
PN	US2003194781-A1.			
PD	16-OCT-2003.			
PA	(GETH) GENENTECH INC.			
Query Match	100.0%;	Score 1688;	DB 7;	Length 321;
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;		
RESULT 36				
ID	ADE16878 standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.			
PN	US2003203433-A1.			
PD	30-OCT-2003.			
PA	(GETH) GENENTECH INC.			
Query Match	100.0%;	Score 1688;	DB 7;	Length 321;
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;		
RESULT 37				
ID	ADF46892 standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.			
PN	US2003195333-A1.			
PD	16-OCT-2003.			
PA	(GETH) GENENTECH INC.			
Query Match	100.0%;	Score 1688;	DB 7;	Length 321;
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;		
RESULT 38				
ID	ADG52649 standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.			
PN	US2003216561-A1.			
PD	20-NOV-2003.			
PA	(GETH) GENENTECH INC.			
Query Match	100.0%;	Score 1688;	DB 7;	Length 321;
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;		
RESULT 39				
ID	ADG59969 standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.			
PN	US2003206915-A1.			
PD	06-NOV-2003.			
PA	(GETH) GENENTECH INC.			
Query Match	100.0%;	Score 1688;	DB 7;	Length 321;
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;		
RESULT 40				
ID	ADH62529 standard; protein; 321 AA.			
DE	Human PRO362 protein encoded by DNA45416.			
PN	US2003171568-A1.			
PD	11-SEP-2003.			
PA	(ASHK/) ASHKENAZI A.			
PA	(FONG/) FONG S.			
PA	(GODD/) GODDARD A.			
PA	(GURN/) GURNEY A L.			
PA	(NAPI/) NAPIER M A.			
PA	(TUMA/) TUMAS D.			
PA	(WOOD/) WOOD W I.			
Query Match	100.0%;	Score 1688;	DB 7;	Length 321;
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;		
RESULT 41				
ID	ADI60729 standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.			
PN	US2003077700-A1.			
PD	24-APR-2003.			
PA	(GETH) GENENTECH INC.			
Query Match	100.0%;	Score 1688;	DB 7;	Length 321;
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;		
RESULT 42				
ID	ADH62529 standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.			
PN	US2003171568-A1.			
PD	11-SEP-2003.			
PA	(ASHK/) ASHKENAZI A.			
PA	(FONG/) FONG S.			
PA	(GODD/) GODDARD A.			
PA	(GURN/) GURNEY A L.			
PA	(NAPI/) NAPIER M A.			
PA	(TUMA/) TUMAS D.			
PA	(WOOD/) WOOD W I.			
Query Match	100.0%;	Score 1688;	DB 7;	Length 321;
Best Local Similarity	100.0%;	Pred. No. 9.4e-136;		
RESULT 43				
ID	ADH89487 standard; protein; 321 AA.			
DE	Human secreted/transmembrane protein, PRO362.			
PN	US2003130181-A1.			
PD	10-JUL-2003.			
PA	(ASHK/) ASHKENAZI A J.			
PA	(BAKE/) BAKER K P.			
PA	(BOTS/) BOTSTEIN D.			
PA	(DESN/) DESNOVERS L.			
PA	(EATO/) EATON D L.			
PA	(FERR/) FERRARA N.			
PA	(FILV/) FILVAROFF E.			
PA	(FONG/) FONG S.			
PA	(GAOW/) GAO W.		</	

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RESULT 48
ID ADF40443 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003199021-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 49
ID ADF23387 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003203402-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 50
ID ADF33370 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003194780-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 51
ID ADF26837 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003199436-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 52
ID ADF27473 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003199437-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 53
ID ADF41067 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003199435-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 54
ID ADF32746 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003211091-A1.
PD 13-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 55
ID ADF25112 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003211092-A1.
PD 13-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 56
ID ADF26213 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003199674-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 57
ID ADF32429 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003194410-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 58
ID ADF46239 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003195344-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 59
ID ADG50225 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003207803-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 60
ID ADG49601 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003215905-A1.
PD 20-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 61
ID ADG51473 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003215908-A1.
PD 20-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 62
ID ADG48977 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003216305-A1.
PD 20-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 63
ID ADG48353 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2003216560-A1.
PD 20-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 64
ID ADG50849 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2004005312-A1.
PD 08-JAN-2004.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 65
ID ADG58793 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2004005657-A1.
PD 08-JAN-2004.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Score 1688; DB 8; Length 321;
RESULT 66
ID ADG62249 standard; protein; 321 AA.
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DE Human secreted/transmembrane protein, PRO362.
PN US2004006219-A1.
PD 08-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 8; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 67
ID ADH25274 standard; protein; 321 AA.
DE Human neurotrophin homologue related protein sequence SEQ ID NO:52.
PN EP1386931-A1.
PD 04-FEB-2004.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 8; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 68
ID ADM17051 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2004048332-A1.
PD 11-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 8; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 69
ID ADL66885 standard; protein; 321 AA.
DE Human secreted/transmembrane protein, PRO362.
PN US2004063921-A1.
PD 01-APR-2004.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 8; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 70
ID ADN35285 standard; protein; 321 AA.
DE Human STIGMA protein.
PN WO2004031105-A2.
PD 15-APR-2004.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 8; Length 321;
Best Local Similarity 100.0%; Pred. No. 9.4e-136;
RESULT 71
ID AAB19396 standard; protein; 399 AA.
DE Amino acid sequence of a human secreted protein.
PN WO200061755-A2.
PD 19-OCT-2000.
PA (CHIR) CHIRON CORP.
Query Match 100.0%; Score 1688; DB 3; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 72
ID AAB04230 standard; protein; 399 AA.
DE Human gene 11 encoded secreted protein HMSOW51, SEQ ID NO:85.
PN WO200136432-A2.
PD 25-MAY-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 4; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 73
ID AAE04208 standard; protein; 399 AA.
DE Human gene 11 encoded secreted protein HMSOW51, SEQ ID NO:63.
PN WO200136432-A2.
PD 25-MAY-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 4; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 74
ID ABG64488 standard; protein; 399 AA.
DE Human albumin fusion protein #1163.
PN WO200177137-A1.
PD 18-OCT-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 5; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 75
ID ABG64487 standard; protein; 399 AA.
DE Human albumin fusion protein #1162.

PN WO200177137-A1.
PD 18-OCT-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 5; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 76
ID ABU03567 standard; protein; 399 AA.
DE Angiogenesis-associated human protein sequence #112.
PN WO200279492-A2.
PD 10-OCT-2002.
PA (EOSB-) EOS BIOTECHNOLOGY INC.
Query Match 100.0%; Score 1688; DB 6; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 77
ID ADA57183 standard; protein; 399 AA.
DE Human secreted protein #466.
PN WO2002102994-A2.
PD 27-DEC-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 6; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 78
ID ADA41052 standard; protein; 399 AA.
DE Human secreted protein.
PN WO2002102993-A2.
PD 27-DEC-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 6; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 79
ID ABR47904 standard; protein; 399 AA.
DE Human secreted protein, SEQ ID 795.
PN WO200295010-A2.
PD 28-NOV-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 6; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 80
ID ABR00157 standard; protein; 399 AA.
DE Human gene 147 encoded secreted protein HSDEK49, SEQ ID NO:446.
PN WO200276488-A1.
PD 03-OCT-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 6; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 81
ID ADB91646 standard; protein; 399 AA.
DE Human secreted protein #SEQ ID 592.
PN WO2003004622-A2.
PD 16-JAN-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 7; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 82
ID ADC74298 standard; protein; 399 AA.
DE Human secreted protein - SEQ ID 931.
PN WO2003038063-A2.
PD 08-MAY-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 100.0%; Score 1688; DB 7; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 83
ID ADL77755 standard; protein; 399 AA.
DE Albumin fusion protein related therapeutic protein X, SEQ ID NO 1237.
PN US2004010134-A1.
PD 15-JAN-2004.
PA (ROSE/) ROSEN C A.
Query Match 100.0%; Score 1688; DB 8; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 84
ID ADL77754 standard; protein; 399 AA.
DE Albumin fusion protein related therapeutic protein X, SEQ ID NO 1236.

PN US2004010134-A1.
PD 15-JAN-2004.
PA (ROSE/) ROSEN C A.
PA (HASE/) HASELTINE W A.
Query Match 100.0%; Score 1688; DB 8; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 85
ID ADL67172 standard; protein; 399 AA.
DE Human B7-H6 protein SEQ ID NO:42.
PN WO2004022594-A2.
PD 18-MAR-2004.
PA (CYTO-) CYTOS BIOTECHNOLOGY AG.
Query Match 100.0%; Score 1688; DB 8; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 86
ID ADN35311 standard; protein; 399 AA.
DE Human STIGMA protein.
PN WO2004031105-A2.
PD 15-APR-2004.
PA (GETH-) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 8; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 87
ID ABM2416 standard; protein; 399 AA.
DE Tumour-associated antigenic target (TAT) polypeptide PRO34043, SEQ:6205.
PN WO2004030615-A2.
PD 15-APR-2004.
PA (GETH-) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 8; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 88
ID ADP23762 standard; protein; 399 AA.
DE PRO polypeptide SEQ ID NO:940.
PN WO2004041170-A2.
PD 21-MAY-2004.
PA (GETH-) GENENTECH INC.
Query Match 100.0%; Score 1688; DB 8; Length 399;
Best Local Similarity 100.0%; Pred. No. 1.3e-135;
RESULT 89
ID AAM3874 standard; protein; 399 AA.
DE Human polypeptide, SEQ ID NO: 3986.
PN EP1130094-A2.
PD 05-SEP-2001.
PA (HELI-) HELIX RES INST.
Query Match 99.5%; Score 1680; DB 4; Length 399;
Best Local Similarity 99.4%; Pred. No. 6.1e-135;
RESULT 90
ID ADL31953 standard; protein; 399 AA.
DE Human protein encoded by a full length cDNA clone SeqID 3986.
PN EP1396543-A2.
PD 10-MAR-2004.
PA (REAS-) RES ASSOC BIOTECHNOLOGY.
Query Match 99.5%; Score 1680; DB 8; Length 399;
Best Local Similarity 99.4%; Pred. No. 6.1e-135;
RESULT 91
ID AAE04290 standard; protein; 386 AA.
DE Human gene 11 encoded secreted protein fragment, SEQ ID NO:154.
PN WO200136432-A2.
PD 25-MAY-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 96.3%; Score 1625; DB 4; Length 386;
Best Local Similarity 100.0%; Pred. No. 3e-130;
RESULT 92
ID ADL67174 standard; protein; 281 AA.
DE Human B7-H6 (ECD) protein SEQ ID NO:44.
PN WO2004022594-A2.
PD 18-MAR-2004.
PA (CYTO-) CYTOS BIOTECHNOLOGY AG.
Query Match 87.7%; Score 1480; DB 8; Length 281;
Best Local Similarity 100.0%; Pred. No. 4.9e-118;
RESULT 93
ID ADA57531 standard; protein; 305 AA.
DE Human secreted protein #466.

PN WO2002102994-A2.
PD 27-DEC-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 67.4%; Score 1137; DB 6; Length 305;
Best Local Similarity 70.4%; Pred. No. 1.2e-88;
RESULT 94
ID ADA41415 standard; protein; 305 AA.
DE Human secreted protein.
PN WO2002102993-A2.
PD 27-DEC-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 67.4%; Score 1137; DB 6; Length 305;
Best Local Similarity 70.4%; Pred. No. 1.2e-88;
RESULT 95
ID ABR48114 standard; protein; 305 AA.
DE Human secreted protein, SEQ ID 1005.
PN WO200295010-A2.
PD 28-NOV-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 67.4%; Score 1137; DB 6; Length 305;
Best Local Similarity 70.4%; Pred. No. 1.2e-88;
RESULT 96
ID ABR00282 standard; protein; 305 AA.
DE Human gene 147 encoded secreted protein HSDEK49, SEQ ID NO:571.
PN WO200276488-A1.
PD 03-OCT-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 67.4%; Score 1137; DB 6; Length 305;
Best Local Similarity 70.4%; Pred. No. 1.2e-88;
RESULT 97
ID ADB91804 standard; protein; 305 AA.
DE Human secreted protein #SEQ ID 750.
PN WO2003004622-A2.
PD 16-JAN-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 67.4%; Score 1137; DB 7; Length 305;
Best Local Similarity 70.4%; Pred. No. 1.2e-88;
RESULT 98
ID ADC74547 standard; protein; 305 AA.
DE Human secreted protein - SEQ ID 1180.
PN WO2003038063-A2.
PD 08-MAY-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 67.4%; Score 1137; DB 7; Length 305;
Best Local Similarity 70.4%; Pred. No. 1.2e-88;
RESULT 99
ID ADN35312 standard; protein; 305 AA.
DE Human short STIGMA protein.
PN WO2004031105-A2.
PD 15-APR-2004.
PA (GETH-) GENENTECH INC.
Query Match 67.4%; Score 1137; DB 8; Length 305;
Best Local Similarity 70.4%; Pred. No. 1.2e-88;
RESULT 100
ID AAY30814 standard; protein; 306 AA.
DE Human secreted protein encoded from gene 4.
PN WO9940100-A1.
PD 12-AUG-1999.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 67.4%; Score 1137; DB 2; Length 306;
Best Local Similarity 70.4%; Pred. No. 1.2e-88;
RESULT 101
ID AAM93588 standard; protein; 184 AA.
DE Human polypeptide, SEQ ID NO: 3387.
PN EP1130094-A2.
PD 05-SEP-2001.
PA (HELI-) HELIX RES INST.
Query Match 58.2%; Score 983; DB 4; Length 184;
Best Local Similarity 100.0%; Pred. No. 8.8e-76;
RESULT 102
ID ADL31354 standard; protein; 184 AA.
DE Human protein encoded by a full length cDNA clone SeqID 3387.
PN EP1396543-A2.

PD 10-MAR-2004.
 PA (REAS-) RES ASSOC BIOTECHNOLOGY.
 Query Match 58.2%; Score 983; DB 8; Length 184;
 Best Local Similarity 100.0%; Pred. No. 8.8e-76;
 RESULT 103
 ID AAY82322 standard; protein; 175 AA.
 DE Human protein transport molecule (PTAM) SEQ ID NO:6.
 PD WO200012703-A2.
 PD 09-MAR-2000.
 PA (INCY-) INCYTE PHARM INC.
 Query Match 51.5%; Score 870; DB 3; Length 175;
 Best Local Similarity 64.7%; Pred. No. 3.8e-66;
 RESULT 104
 ID ADK70486 standard; protein; 175 AA.
 DE Respiratory disease differentially expressed protein #52.
 PD WO2003101283-A2.
 PD 11-DEC-2003.
 PA (INCY-) INCYTE CORP.
 Query Match 51.5%; Score 870; DB 8; Length 175;
 Best Local Similarity 64.7%; Pred. No. 3.8e-66;
 RESULT 105
 ID ADL67144 standard; protein; 280 AA.
 DE Mouse B7-H6 protein SEQ ID NO:14.
 PD WO2004022594-A2.
 PD 18-MAR-2004.
 PA (CYTO-) CYTOS BIOTECHNOLOGY AG.
 Query Match 43.1%; Score 728; DB 8; Length 280;
 Best Local Similarity 47.0%; Pred. No. 1e-53;
 RESULT 106
 ID ADN5313 standard; protein; 280 AA.
 DE Murine STGMA protein.
 PD WO2004031105-A2.
 PD 15-APR-2004.
 PA (GETH-) GENENTECH INC.
 Query Match 43.1%; Score 728; DB 8; Length 280;
 Best Local Similarity 47.0%; Pred. No. 1e-53;
 RESULT 107
 ID ADL67146 standard; protein; 188 AA.
 DE Mouse B7-H6 (ECD) protein SEQ ID NO:16.
 PD WO2004022594-A2.
 PD 18-MAR-2004.
 PA (CYTO-) CYTOS BIOTECHNOLOGY AG.
 Query Match 36.3%; Score 613; DB 8; Length 188;
 Best Local Similarity 45.7%; Pred. No. 4e-44;
 RESULT 108
 ID AAE10596 standard; protein; 93 AA.
 DE Human macrophage-expressed protein #21.
 PD WO200164839-A2.
 PD 07-SEP-2001.
 PA (HYSE-) HYSEQ INC.
 Query Match 29.1%; Score 492; DB 4; Length 93;
 Best Local Similarity 98.9%; Pred. No. 3.5e-34;
 RESULT 109
 ID ADH80723 standard; protein; 300 AA.
 DE Human polypeptide #40.
 PD US2003232054-A1.
 PD 18-DEC-2003.
 PA (TANG/) TANG Y T.
 PA (LIUC/) LIU C.
 PA (ASUN/) ASUNDI V.
 PA (CHEN/) CHEN R.
 PA (QIAN/) QIAN X B.
 PA (WANG/) WANG Z W.
 PA (WEHR/) WEHRMAN T.
 PA (ZHAN/) ZHANG J.
 PA (ZHOU/) ZHOU P.
 PA (CAO/) CAO Y.
 PA (DRMA/) DRMANAC R T.
 Query Match 10.7%; Score 180; DB 8; Length 300;
 Best Local Similarity 27.9%; Pred. No. 8.2e-07;
 RESULT 110
 ID AAY23321 standard; protein; 299 AA.
 DE Amino acid sequence of the PRO301 polypeptide.

PN WO9927098-A2.
 PD 03-JUN-1999.
 PA (GETH-) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 2; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 111
 ID AAW74464 standard; protein; 299 AA.
 DE F11 antigen protein sequence.
 PN WO9902561-A1.
 PD 21-JAN-1999.
 PA (SMIK-) SMITHKLINE BEECHAM CORP.
 Query Match 10.6%; Score 178.5; DB 2; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 112
 ID AAY08071 standard; protein; 299 AA.
 DE Human PRO307 protein.
 PN WO9914241-A2.
 PD 25-MAR-1999.
 PA (GETH-) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 2; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 113
 ID AAV13364 standard; protein; 299 AA.
 DE Amino acid sequence of protein PRO301.
 PN WO9914328-A2.
 PD 25-MAR-1999.
 PA (GETH-) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 2; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 114
 ID AAY76011 standard; protein; 299 AA.
 DE Human A33 receptor homologue, SEQ ID NO:189.
 PN WO9955865-A1.
 PD 04-NOV-1999.
 PA (GENE-) GENESIS RES & DEV CORP LTD.
 Query Match 10.6%; Score 178.5; DB 3; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 115
 ID AAY76076 standard; protein; 299 AA.
 DE Human A33 receptor homologue, SEQ ID NO:331.
 PN WO9955865-A1.
 PD 04-NOV-1999.
 PA (GENE-) GENESIS RES & DEV CORP LTD.
 Query Match 10.6%; Score 178.5; DB 3; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 116
 ID AAY70670 standard; protein; 299 AA.
 DE Human PRO301 protein.
 PN WO200015797-A2.
 PD 23-MAR-2000.
 PA (GETH-) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 3; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 117
 ID AAB24405 standard; protein; 299 AA.
 DE Human PRO301 protein sequence SEQ ID NO:90.
 PN WO200032221-A2.
 PD 08-JUN-2000.
 PA (GETH-) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 3; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 118
 ID AAY95344 standard; protein; 299 AA.
 DE Human PRO301 antitumour protein.
 PN WO200037638-A2.
 PD 29-JUN-2000.
 PA (GETH-) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 3; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 119
 ID AAB80232 standard; protein; 299 AA.
 DE Human PRO301 protein.
 PN WO200104311-A1.

PD 18-JAN-2001.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 4; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 120
 ID AAM93577 standard; protein; 299 AA.
 DE Human polypeptide, SEQ ID NO: 3365.
 PN EPI130094-A2.
 PD 05-SEP-2001.
 PA (HELI-) HELIX RES INST.
 Query Match 10.6%; Score 178.5; DB 4; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 121
 ID AAB56015 standard; protein; 299 AA.
 DE Skin cell protein, SEQ ID NO: 331.
 PN WO200069884-A2.
 PD 23-NOV-2000.
 PA (GENE-) GENESIS RES & DEV CORP LTD.
 Query Match 10.6%; Score 178.5; DB 4; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 122
 ID AAB55950 standard; protein; 299 AA.
 DE Skin cell protein, SEQ ID NO: 189.
 PN WO200069884-A2.
 PD 23-NOV-2000.
 PA (GENE-) GENESIS RES & DEV CORP LTD.
 Query Match 10.6%; Score 178.5; DB 4; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 123
 ID AAB31202 standard; protein; 299 AA.
 DE Amino acid sequence of human polypeptide PRO301.
 PN WO20007037-A2.
 PD 21-DEC-2000.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 4; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 124
 ID AAU00823 standard; protein; 299 AA.
 DE Human immune response protein PRO301 (UNQ264).
 PN WO200119991-A1.
 PD 22-MAR-2001.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 4; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 125
 ID AAU12354 standard; protein; 299 AA.
 DE Human PRO301 polypeptide sequence.
 PN WO200140466-A2.
 PD 07-JUN-2001.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 4; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 126
 ID AAB53086 standard; protein; 299 AA.
 DE Human angiogenesis-associated protein PRO301, SEQ ID NO:119.
 PN WO200053753-A2.
 PD 14-SEP-2000.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 4; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 127
 ID AAU14405 standard; protein; 299 AA.
 DE Human novel protein #276.
 PN WO200155437-A2.
 PD 02-AUG-2001.
 PA (HYSE-) HYSEQ INC.
 Query Match 10.6%; Score 178.5; DB 4; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 128
 ID AAU14404 standard; protein; 299 AA.
 DE Human novel protein #275.
 PN WO200155437-A2.
 PD 02-AUG-2001.

PA (HYSE-) HYSEQ INC.
 Query Match 10.6%; Score 178.5; DB 4; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 129
 ID AAU14168 standard; protein; 299 AA.
 DE Human novel protein #39.
 PN WO200155437-A2.
 PD 02-AUG-2001.
 PA (HYSE-) HYSEQ INC.
 Query Match 10.6%; Score 178.5; DB 4; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 130
 ID AAE03896 standard; protein; 299 AA.
 DE Human gene 23 encoded secreted protein fragment, SEQ ID NO:148.
 PN WO200136440-A1.
 PD 25-MAY-2001.
 PA (HUMA-) HUMAN GENOME SCI INC.
 Query Match 10.6%; Score 178.5; DB 4; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 131
 ID AAE03840 standard; protein; 299 AA.
 DE Human gene 23 encoded secreted protein HACAA29, SEQ ID NO: 86.
 PN WO200136440-A1.
 PD 25-MAY-2001.
 PA (HUMA-) HUMAN GENOME SCI INC.
 Query Match 10.6%; Score 178.5; DB 4; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 132
 ID AAE03870 standard; protein; 299 AA.
 DE Human gene 23 encoded secreted protein HACAA29, SEQ ID NO:116.
 PN WO200136440-A1.
 PD 25-MAY-2001.
 PA (HUMA-) HUMAN GENOME SCI INC.
 Query Match 10.6%; Score 178.5; DB 4; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 133
 ID ABB90290 standard; protein; 299 AA.
 DE Human polypeptide SEQ ID NO 2666.
 PN WO200190304-A2.
 PD 29-NOV-2001.
 PA (HUMA-) HUMAN GENOME SCI INC.
 Query Match 10.6%; Score 178.5; DB 5; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 134
 ID ABB84843 standard; protein; 299 AA.
 DE Human PRO301 protein sequence SEQ ID NO:54.
 PN WO200200690-A2.
 PD 03-JAN-2002.
 PA (GETH) GENENTECH INC.
 Query Match 10.6%; Score 178.5; DB 5; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 135
 ID ABG64551 standard; protein; 299 AA.
 DE Human albumin fusion protein #1226.
 PN WO200177137-A1.
 PD 18-OCT-2001.
 PA (HUMA-) HUMAN GENOME SCI INC.
 Query Match 10.6%; Score 178.5; DB 5; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 136
 ID ABG64552 standard; protein; 299 AA.
 DE Human albumin fusion protein #1227.
 PN WO200177137-A1.
 PD 18-OCT-2001.
 PA (HUMA-) HUMAN GENOME SCI INC.
 Query Match 10.6%; Score 178.5; DB 5; Length 299;
 Best Local Similarity 26.5%; Pred. No. 1.1e-06;
 RESULT 137
 ID ABB72215 standard; protein; 299 AA.
 DE Human protein isolated from skin cells SEQ ID NO: 331.
 PN WO200190357-A1.
 PD 29-NOV-2001.
 PA (GENE-) GENESIS RES & DEV CORP LTD.

Query Match 10.6%; Score 178.5; DB 5; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 138
ID ABB72150 standard; protein; 299 AA.
DE Human protein isolated from skin cells SEQ ID NO: 189.
PN WO200190357-A1.
PD 29-NOV-2001.
PA (GENE-) GENESIS RES & DEV CORP LTD.
Query Match 10.6%; Score 178.5; DB 5; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 139
ID ABB95449 standard; protein; 299 AA.
DE Human angiogenesis related protein PRO301 SEQ ID NO: 54.
PN WO200208284-A2.
PD 31-JAN-2002.
PA (GETH) GENENTECH INC.
PA (BAKE/) BAKER K P.
PA (FERR/) FERRARA N.
PA (GERB/) GERBER H.
PA (GERR/) GERRITSEN M E.
PA (GODD/) GODDARD A.
PA (GODO/) GODOWSKI P J.
PA (GURN/) GURNEY A L.
PA (HILL/) HILLAN K J.
PA (MARS/) MARSTERS S A.
PA (PANU/) PAN J.
PA (PAON/) PAONI N F.
PA (STEP/) STEPHAN J F.
PA (WATA/) WATANABE C K.
PA (WILL/) WILLIAMS P M.
PA (WOOD/) WOOD W I.
Query Match 10.6%; Score 178.5; DB 5; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 140
ID ABU71610 standard; protein; 299 AA.
DE Human PRO polypeptide #21.
PN US2002146709-A1.
PD 10-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 141
ID ABO17798 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003032156-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 142
ID ABU71465 standard; protein; 299 AA.
DE Human PRO polypeptide #21.
PN US2002192659-A1.
PD 19-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 143
ID ABO25173 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003040014-A1.
PD 27-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 144
ID ABU81052 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003004311-A1.
PD 02-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;

RESULT 145
ID ABU71911 standard; protein; 299 AA.
DE Human secreted/transmembrane protein PRO301.
PN US2003003530-A1.
PD 02-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 146
ID ABO01794 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2002197671-A1.
PD 26-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 147
ID ABU66752 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003036180-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 148
ID ABU54367 standard; protein; 299 AA.
DE Human secreted/transmembrane protein PRO301.
PN US2002132240-A1.
PD 19-SEP-2002.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 149
ID ABU67291 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003032063-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 150
ID ABO47382 standard; protein; 299 AA.
DE Human secreted/transmembrane polypeptide PRO301.
PN US2003044839-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 151
ID ABU59833 standard; protein; 299 AA.
DE Novel secreted and transmembrane protein PRO301.
PN US2003017563-A1.
PD 23-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 152
ID ABO25023 standard; protein; 299 AA.
DE Human secreted/transmembrane protein (PRO) #183.
PN US2003036179-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 153
ID ABU64519 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #23.
PN US2002160374-A1.
PD 31-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 154

ID ABU72059 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2002177165-A1.
PD 28-NOV-2002.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 155
ID ABU67365 standard; protein; 299 AA.
DE Human secreted protein PRO301.
PN US2003023054-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 156
ID ABU67160 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003032062-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 157
ID ABO14895 standard; protein; 299 AA.
DE Human secreted / transmembrane polypeptide PRO301.
PN US2003036060-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 158
ID ABU07736 standard; protein; 299 AA.
DE Human A-33 related antigen PRO301.
PN US2002182206-A1.
PD 05-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 159
ID AAO16451 standard; protein; 299 AA.
DE Human junctional adhesion molecule 1 (hujam1).
PN WO2003008541-A2.
PD 30-JAN-2003.
PA (ELIL) LILLY & CO ELI.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 160
ID ABU67028 standard; protein; 299 AA.
DE Human secreted/transmembrane, PRO, protein SEQ ID 366.
PN US2003032155-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 161
ID ABU69642 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003017463-A1.
PD 23-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 162
ID ABU79802 standard; protein; 299 AA.
DE Human secreted/transmembrane protein PRO301.
PN US2003032057-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 163
ID ABO14824 standard; protein; 299 AA.

DE Human secreted / transmembrane polypeptide PRO301.
PN US2003027143-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 164
ID ADA45985 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003022328-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 165
ID ADA76316 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003073212-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 166
ID ADB29324 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003092002-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 167
ID ADA18966 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003054517-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 168
ID ADA61589 standard; protein; 299 AA.
DE Homo sapiens.
PN US2003049816-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 169
ID ADB19374 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003068796-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 170
ID ADB27915 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082704-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 171
ID ADA86394 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082711-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 172
ID ADB15958 standard; protein; 299 AA.
DE Human PRO polypeptide #183.

PN US2003087350-A1.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 173
ID ADA47744 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003073215-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 174
ID ADA18180 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003039971-A1.
PD 27-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 175
ID ABO32776 standard; protein; 299 AA.
DE Human secreted/transmembrane protein PRO301.
PN US2003045693-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 176
ID ADA67539 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003068795-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 177
ID ADB30546 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003068794-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 178
ID ADA85842 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082693-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 179
ID ADA97054 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082705-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 180
ID ADA79358 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082763-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 181
ID ADA87497 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087345-A1.

PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 182
ID ADB16699 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003087349-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 183
ID ABO34836 standard; protein; 299 AA.
DE Human PRO polypeptide #21.
PN US2003044793-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 184
ID ADA16155 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003049621-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 185
ID ADA91791 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082694-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 186
ID ADB14854 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003087351-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 187
ID ADA47263 standard; protein; 299 AA.
DE Human secreted/transmembrane polypeptide PRO301.
PN US2003044844-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 188
ID ADB18815 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003073211-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 189
ID ADA94030 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003077722-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 190
ID ADB19926 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082691-A1.
PD 01-MAY-2003.

PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 191
ID ADB13238 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082710-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 192
ID ABO43331 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003044945-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 193
ID ADA74492 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003068798-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 194
ID ADA42300 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003054401-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 195
ID ADB24725 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US2003077713-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 196
ID ADA82249 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082701-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 197
ID ADA75212 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003073216-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 198
ID ADA85290 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082695-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 199
ID ADA84738 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082708-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 200
ID ABO17514 standard; protein; 299 AA.
DE Human PRO polypeptide #21.
PN US2003064367-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 201
ID ADB29994 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003073214-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 202
ID ADA80522 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082761-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 203
ID ADA75764 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082703-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 204
ID ADA46989 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003073210-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 205
ID ADB25285 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US2003077715-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 206
ID ADA93461 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003077721-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 207
ID ADB26811 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003092147-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 208
ID ADB31098 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003096386-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;


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Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 209
ID ADA61026 standard; protein; 299 AA.
DE Homo sapiens.
PN US2003049817-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 210
ID ADB24173 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US2003077714-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 211
ID ADA96502 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082690-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 212
ID ADA81074 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082702-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 213
ID ADA95950 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082759-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 214
ID ADB26259 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082760-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 215
ID ADB21744 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082765-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 6; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 216
ID ADA77523 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003068797-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 217
ID ADB18263 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US200307710-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 218
ID ADA86946 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082709-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 219
ID ADA16579 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003039969-A1.
PD 27-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 220
ID ADA13008 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003049622-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 221
ID ADA41876 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003082540-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 222
ID ADA88049 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082700-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 223
ID ADA46437 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003054516-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 224
ID ADA17223 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003017498-A1.
PD 23-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 225
ID ADA42726 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003054351-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 226
ID ADB28467 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082699-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 227
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ID ADB29019 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082706-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 228
ID ADA76971 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003059909-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 229
ID ADA88601 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003073213-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 230
ID ADA97606 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082686-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 231
ID ADB27363 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003022239-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 232
ID ADB22296 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087344-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 233
ID ABO19860 standard; protein; 299 AA.
DE Human secreted/transmembrane protein PRO302.
PN US2003044902-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 234
ID ABO17575 standard; protein; 299 AA.
DE Human PRO polypeptide #21.
PN US2003064923-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 235
ID ADA66987 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003068793-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 236
ID ADB22848 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003077711-A1.

PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 237
ID ADB23621 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US2003077712-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 238
ID ADA92343 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082712-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 239
ID ADB15406 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003087352-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 240
ID ADB38658 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082766-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 241
ID ADB38106 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087347-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 242
ID ADB66578 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082689-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 243
ID ADB89658 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082698-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 244
ID ADB90390 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082762-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 245
ID ADB77645 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003077654-A1.
PD 24-APR-2003.

PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 246
ID ADB39491 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082764-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 247
ID ADB47481 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003082542-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 248
ID ADB47114 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082687-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 249
ID ADB6721 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003082697-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 250
ID ADB77326 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082696-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 251
ID ADB34483 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US200307717-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 252
ID ADB35587 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US200307719-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 253
ID ADB33931 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US200307716-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 254
ID ADB35035 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US200307718-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.

Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 255
ID ADB36139 standard; protein; 299 AA.
DE Human PRO polypeptide SEQ ID NO 366.
PN US200307720-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 256
ID ADB46534 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082692-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 257
ID ADC28427 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003059772-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 258
ID ADC39627 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003059828-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 259
ID ADC40141 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003059829-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 260
ID ADC18969 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003036061-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 261
ID ADC34265 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003036094-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 262
ID ADC29320 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003049676-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 263
ID ADC28851 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003049677-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;

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Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 264
ID ADC40736 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein, #25.
PN US2003054400-A1.
PD 20-MAR-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 265
ID ADC19393 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003054441-A1.
PD 20-MAR-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 266
ID ADC33841 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003073077-A1.
PD 17-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 267
ID ADC12911 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003073079-A1.
PD 17-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 268
ID ADC50407 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003092106-A1.
PD 15-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 269
ID ADC71954 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003092107-A1.
PD 15-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 270
ID ADC59933 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003092105-A1.
PD 15-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 271
ID ADC52940 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein Seq ID366.
PN US2003087365-A1.
PD 08-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 272
ID ADC57294 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein Seq ID366.
PN US2003087366-A1.
PD 08-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 273
ID ADC57294 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein, #25.
PN US2003087366-A1.
PD 08-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 274
ID ADC5960 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087361-A1.
PD 08-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 275
ID ADC65487 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003087362-A1.
PD 08-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 276
ID ADC54585 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein Seq ID366.
PN US2003087363-A1.
PD 08-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 277
ID ADC53546 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein Seq ID366.
PN US2003087364-A1.
PD 08-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 278
ID ADC59069 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein Seq ID366.
PN US2003087359-A1.
PD 08-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 279
ID ADC55947 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein Seq ID366.
PN US2003087360-A1.
PD 08-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 280
ID ADC58517 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein Seq ID366.
PN US2003087346-A1.
PD 08-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 281
ID ADC12363 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003082541-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 282
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ID ADD03191 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003092104-A1.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 283
ID ADC90183 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087348-A1.
PD 08-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 284
ID ADC69602 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194773-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 285
ID ADC48491 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194773-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 286
ID ADD10020 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194776-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 287
ID ADD04595 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087334-A1.
PD 08-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 288
ID ADC80551 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003092103-A1.
PD 15-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 289
ID ADD11058 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194774-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 290
ID ADD10343 standard; protein; 299 AA.
DE Human secreted/transmembrane PRO polypeptide #27.
PN US2003105011-A1.
PD 05-JUN-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 291
ID ADC47939 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194771-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 292
ID ADD04918 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003104469-A1.
PD 05-JUN-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 293
ID ADC79999 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087358-A1.
PD 08-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 294
ID ADD11303 standard; protein; 299 AA.
DE Human secreted/transmembrane PRO polypeptide #27.
PN US2003105013-A1.
PD 05-JUN-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 295
ID ADD09468 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194775-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 296
ID ADD03924 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003104381-A1.
PD 05-JUN-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 297
ID ADD03500 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003108983-A1.
PD 12-JUN-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 298
ID ADD4181 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003203438-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 299
ID ADD52320 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194769-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 300
ID ADD53060 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
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PN US2003194792-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 301
ID ADD531612 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003203437-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 302
ID ADD37096 standard; protein; 299 AA.
DE Human secreted/transmembrane PRO polypeptide #27.
PN US2003105012-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 303
ID ADD51768 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194779-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 304
ID ADD02567 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003203431-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 305
ID ADD02001 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003203430-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 306
ID ADD54183 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003203432-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 307
ID ADD92500 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199030-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 308
ID ADD91396 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199055-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 309
ID ADE04010 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199057-A1.

PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 310
ID ADE32307 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003194765-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 311
ID ADE22239 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199056-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 312
ID ADD79463 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003203428-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 313
ID ADE41999 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194772-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 314
ID ADE17816 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199023-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 315
ID ADD91948 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199053-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 316
ID ADE33411 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003194767-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 317
ID ADE33963 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003194791-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 318
ID ADD80015 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207417-A1.
PD 06-NOV-2003.

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PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 319
ID ADE42551 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199032-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 320
ID ADE19472 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199025-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 321
ID ADE18920 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199026-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 322
ID ADE18920 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199026-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 323
ID ADE3116 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199033-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 324
ID ADD95905 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199059-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 325
ID ADE22791 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199064-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 326
ID ADD78909 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003203429-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 327
ID ADE32859 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003194766-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 328
ID ADE42551 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199032-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 329
ID ADD80567 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207418-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 330
ID ADD89595 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199028-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 331
ID ADE40879 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199031-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 332
ID ADE04678 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199034-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 333
ID ADE92807 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194777-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 334
ID ADG21516 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207355-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 335
ID ADG23157 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207384-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 336
ID ADF97492 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207370-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 337
ID ADE32859 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003194766-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
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Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 337
ID ADG80556 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207373-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 338
ID ADG80004 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207372-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 339
ID ADG63772 standard; protein; 299 AA.
DE Human secreted/transmembrane polypeptide PRO301.
PN US2003170721-A1.
PD 11-SEP-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 340
ID ADH62528 standard; protein; 299 AA.
DE Human PRO301 protein.
PN US2003171568-A1.
PD 11-SEP-2003.
PA (ASHK/) ASHKENAZI A.
PA (FONG/) FONG S.
PA (GODD/) GODDARD A.
PA (GURN/) GURNEY A L.
PA (NAPI/) NAPIER M A.
PA (TUNA/) TUNAS D.
PA (WOOD/) WOOD W I.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 341
ID ADH59235 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003039972-A1.
PD 27-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 342
ID ADH55296 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207381-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 343
ID ADH55848 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207379-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 344
ID ADI38014 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003054352-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 345
ID ADI64067 standard; protein; 299 AA.

DE Novel human secreted and transmembrane protein PRO301.
PN US2003207385-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 346
ID ADI65016 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207386-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 347
ID ADI63515 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207387-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 348
ID ADH81929 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207388-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 349
ID ADH81377 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207377-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 350
ID ADJ26282 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003054349-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 351
ID ADM82546 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087355-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 352
ID ADN15945 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087353-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 353
ID ADN16574 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087385-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 354
ID ADN15393 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.

PN US2003087356-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 355
ID ADN14841 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003087357-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 7; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 356
ID ADC81103 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003092115-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 357
ID ADE79197 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003135025-A1.
PD 17-JUL-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 358
ID ADD76551 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003100087-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 359
ID ADD87915 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003092113-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 360
ID ADD86319 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003203440-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 361
ID ADE79621 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003130489-A1.
PD 10-JUL-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 362
ID ADE75767 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003211571-A1.
PD 13-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 363
ID ADE73297 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003129592-A1.

PD 10-JUL-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 364
ID AD841304 standard; protein; 299 AA.
DE Human secreted/transmembrane PRO polypeptide #27.
PN US2003100497-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 365
ID ADE23343 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003092108-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 366
ID ADE23895 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003092110-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 367
ID ADE24538 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003092111-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 368
ID ADD87363 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003203439-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 369
ID AD889229 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199062-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 370
ID AD841186 standard; protein; 299 AA.
DE Human secreted/transmembrane polypeptide PRO301.
PN US2003104558-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 371
ID ADE73832 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003148370-A1.
PD 07-AUG-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 372
ID ADE18368 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194794-A1.
PD 16-OCT-2003.

PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 373
ID ADE88677 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199054-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 374
ID ADE99386 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003211576-A1.
PD 13-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 375
ID ADE94697 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199027-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 376
ID ADE91108 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199061-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 377
ID ADE95249 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199052-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 378
ID ADE93359 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199060-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 379
ID ADF34940 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199029-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 380
ID ADE98505 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003211569-A1.
PD 13-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 381
ID ADE92255 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003199051-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.

Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 382
ID ADE90556 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003199063-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 383
ID ADE91703 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003199058-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 384
ID ADE98932 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003211568-A1.
PD 13-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 385
ID ADG40402 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US200325253-A1.
PD 04-DEC-2003.
PA (DESN/) DESNOYERS L.
PA (GODD/) GODDARD A. J.
PA (GODO/) GODOWSKI P. J.
PA (GURN/) GURNEY A L.
PA (MATH/) MATHER J P.
PA (WILL/) WILLIAMS P M.
PA (WOOD/) WOOD W I.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 386
ID ADF73796 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003180312-A1.
PD 25-SEP-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 387
ID ADG02282 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207352-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 388
ID ADG22068 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207360-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 389
ID ADG20138 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207376-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 390
ID ADE92255 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003199051-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.

ID ADF98044 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207422-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 391
ID ADG24261 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207426-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 392
ID ADF98615 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003208055-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 393
ID ADG03446 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207351-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 394
ID ADF99167 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207353-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 395
ID ADG16752 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207359-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 396
ID ADG05211 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207375-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 397
ID ADG19478 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207425-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 398
ID ADF73372 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003166051-A1.
PD 04-SEP-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 399
ID ADG13315 standard; protein; 299 AA.

DE Human PRO polypeptide #183.
PN US2003207357-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 400
ID ADG08372 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207424-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 401
ID ADG15542 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003219885-A1.
PD 27-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 402
ID ADF96940 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207371-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 403
ID ADG06125 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207374-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 404
ID ADG23709 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207389-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 405
ID ADG03998 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207423-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 406
ID ADG24899 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207427-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 407
ID ADG07196 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207350-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 408
ID ADG07748 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.

PN US2003207356-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 409
ID ADG55243 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003194778-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 410
ID ADG60907 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207390-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 411
ID ADG62011 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207428-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 412
ID ADG92215 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003027145-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 413
ID ADG82212 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207358-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 414
ID ADG57451 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207362-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 415
ID ADG56899 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207364-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 416
ID ADG55795 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207365-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 417
ID ADG58555 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207368-A1.

PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 418
ID ADG70921 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207420-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 419
ID ADG92642 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003027146-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 420
ID ADG58003 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207363-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 421
ID ADG53587 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207415-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 422
ID ADG71473 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207421-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 423
ID ADG81660 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207805-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 424
ID ADH30622 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003077723-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 425
ID ADG63621 standard; protein; 299 AA.
DE Human secreted/transmembrane polypeptide PRO301.
PN US2003180796-A1.
PD 25-SEP-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 426
ID ADH11989 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207419-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.

Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 427
ID ADG52411 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207414-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 428
ID ADG54139 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207416-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 429
ID ADG81108 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003194793-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 430
ID ADG56347 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207366-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 431
ID ADH21613 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207378-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 432
ID ADG61459 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207429-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 433
ID ADH28546 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003022331-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 434
ID ADG54691 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207367-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 435
ID ADG59731 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207369-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 436
ID ADH20431 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2004005553-A1.
PD 08-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 437
ID ADH43487 standard; protein; 299 AA.
DE Human PRO polypeptide #27.
PN US2003224984-A1.
PD 04-DEC-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 438
ID ADH07286 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2004006211-A1.
PD 08-JAN-2004.
PA (DESN/) DESNOYERS L.
PA (GODD/) GODDARD A. J.
PA (GODO/) GODOWSKI P. J.
PA (GURN/) GURNEY A. L.
PA (MATH/) MATHER J. P.
PA (WILL/) WILLIAMS P. M.
PA (WOOD/) WOOD W. I.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 439
ID ADH59831 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003215904-A1.
PD 20-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 440
ID ADH06859 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2004005665-A1.
PD 08-JAN-2004.
PA (DESN/) DESNOYERS L.
PA (GODD/) GODDARD A. J.
PA (GODO/) GODOWSKI P. J.
PA (GURN/) GURNEY A. L.
PA (MATH/) MATHER J. P.
PA (WILL/) WILLIAMS P. M.
PA (WOOD/) WOOD W. I.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 441
ID ADI18155 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207361-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 442
ID ADI18601 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003152999-A1.
PD 14-AUG-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 443
ID ADI65321 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003148419-A1.

Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 436
ID ADH20431 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2004005553-A1.
PD 08-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 437
ID ADH43487 standard; protein; 299 AA.
DE Human PRO polypeptide #27.
PN US2003224984-A1.
PD 04-DEC-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 438
ID ADH07286 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2004006211-A1.
PD 08-JAN-2004.
PA (DESN/) DESNOYERS L.
PA (GODD/) GODDARD A. J.
PA (GODO/) GODOWSKI P. J.
PA (GURN/) GURNEY A. L.
PA (MATH/) MATHER J. P.
PA (WILL/) WILLIAMS P. M.
PA (WOOD/) WOOD W. I.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 439
ID ADH59831 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003215904-A1.
PD 20-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 440
ID ADH06859 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2004005665-A1.
PD 08-JAN-2004.
PA (DESN/) DESNOYERS L.
PA (GODD/) GODDARD A. J.
PA (GODO/) GODOWSKI P. J.
PA (GURN/) GURNEY A. L.
PA (MATH/) MATHER J. P.
PA (WILL/) WILLIAMS P. M.
PA (WOOD/) WOOD W. I.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 441
ID ADI18155 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003207361-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 442
ID ADI18601 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003152999-A1.
PD 14-AUG-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 443
ID ADI65321 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003148419-A1.

PD 07-AUG-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 444
ID ADI37584 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003096340-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 445
ID ADG09898 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2004009548-A1.
PD 15-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 446
ID ADH97380 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003190610-A1.
PD 09-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 447
ID ADI15369 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207382-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 448
ID ADG09246 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2004009547-A1.
PD 15-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 449
ID ADI65748 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003148371-A1.
PD 07-AUG-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 450
ID ADI14701 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207383-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 451
ID ADH60491 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2004023331-A1.
PD 05-FEB-2004.
PA (DESN) DESNOYERS L.
PA (GODD) GODDARD A.
PA (GODO) GODOWSKI P J.
PA (GUEN) GUERNEY A L.
PA (MATH) MATHER J P.
PA (WILL) WILLIAMS P M.
PA (WOOD) WOOD W I.
Query Match 10.6%; Score 178.5; DB 8; Length 299;

Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 452
ID ADI18296 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207349-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 453
ID ADJ99548 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003187238-A1.
PD 02-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 454
ID ADL08741 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003186358-A1.
PD 02-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 455
ID ADI47176 standard; protein; 299 AA.
DE Human JAM-1 protein sequence.
PN WO2004003145-A2.
PD 08-JAN-2004.
PA (NAST) NASTECH PHARM CO INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 456
ID ADM25082 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003096233-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 457
ID ADK40844 standard; protein; 299 AA.
DE Human platelet F11 receptor #1.
PN US6699688-B1.
PD 02-MAR-2004.
PA (UYN) UNIV NEW YORK STATE RES FOUND.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 458
ID ADJ63577 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2004039164-A1.
PD 26-FEB-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 459
ID ADM29832 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003190611-A1.
PD 09-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 460
ID ADL77818 standard; protein; 299 AA.
DE Albumin fusion protein related therapeutic protein X, SEQ ID No 1300.
PN US2004010134-A1.
PD 15-JAN-2004.
PA (ROSE) ROSEN C A.
PA (HASE) HASELTINE W A.
Query Match 10.6%; Score 178.5; DB 8; Length 299;

Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 461
ID ADL77819 standard; protein; 299 AA.
DE Albumin fusion protein related therapeutic protein X, SEQ ID No 1301.
PN US2004010134-A1.
PD 15-JAN-2004.
PA (ROSE//) ROSEN C A.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 462
ID ADJ77472 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2004038336-A1.
PD 26-FEB-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 463
ID ADK82832 standard; protein; 299 AA.
DE Human PRO polypeptide #27.
PN US2004043927-A1.
PD 04-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 464
ID ADJ65594 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2004038335-A1.
PD 26-FEB-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 465
ID ADL31332 standard; protein; 299 AA.
DE Human protein encoded by a full length cDNA clone SeqID 3365.
PN EPI396543-A2.
PD 10-MAR-2004.
PA (REAS-) RES ASSOC BIOTECHNOLOGY.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 466
ID ADM27730 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2004048333-A1.
PD 11-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 467
ID ADL26800 standard; protein; 299 AA.
DE Human JAM1 protein SEQ ID NO:54.
PN WO2004022778-A1.
PD 18-MAR-2004.
PA (GARV-) GARVAN INST MEDICAL RES.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 468
ID ADM42454 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2004058424-A1.
PD 25-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 469
ID ADO06154 standard; protein; 299 AA.
DE Human PRO polypeptide #21.
PN US6686451-B1.
PD 03-FEB-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 470
ID ADN35284 standard; protein; 299 AA.
DE Human PRO301 protein.
PN WO2004031105-A2.
PD 15-APR-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 471
ID ADN05140 standard; protein; 299 AA.
DE Antipeptidic protein sequence #749.
PN WO2004028479-A2.
PD 08-APR-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 472
ID ADM28316 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2004077064-A1.
PD 22-APR-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 473
ID ADQ95890 standard; protein; 299 AA.
DE T cell activation associated protein #34.
PN WO2004058805-A2.
PD 15-JUL-2004.
PA (ASAH-) ASahi KASEI PHARMA CORP.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 474
ID ADL11006 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2004137561-A1.
PD 15-JUL-2004.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 475
ID ADL17915 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2004147017-A1.
PD 29-JUL-2004.
PA (ASHK/) ASHENAZI A.
PA (BOTS/) BOTSTEIN D.
PA (DESN/) DESNOYERS L.
PA (EATO/) EATON D L.
PA (FERR/) FERRARA N.
PA (FILV/) FILVAROFF E.
PA (FONG/) FONG S.
PA (GAOW/) GAO W.
PA (GERB/) GERBER H.
PA (GERR/) GERRITSEN M E.
PA (GODD/) GODDARD A.
PA (GODO/) GODOWSKI P J.
PA (GRIM/) GRIMALDI C J.
PA (GURN/) GURNEY A L.
PA (HILL/) HILLAN K J.
PA (KLJA/) KLJAVIN I J.
PA (MATH/) MATHER J P.
PA (PANJ/) PAN J.
PA (PAON/) PAONI N F.
PA (ROIM/) ROY M A.
PA (STEW/) STEWART T A.
PA (TUNA/) TUMAS D.
PA (WILL/) WILLIAMS P M.
PA (WOOD/) WOOD W I.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 476

ID ADR27641 standard; protein; 299 AA.
DE Human F11 receptor protein Seq 7.
PN WO2004063327-A2.
PD 29-JUL-2004.
PA (KORN//) KORNECKI E.
PA (BABI//) BABINSKA A.
PA (EHLR//) EHRLICH Y H.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 477
ID ADI95798 standard; protein; 299 AA.
DE Human PRO polypeptide #183.
PN US2003077659-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 478
ID ADI96350 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207354-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 479
ID ADR46577 standard; protein; 299 AA.
DE Human JAM-1, F11 receptor (F11R) transcript variant 4, SEQ ID 8.
PN JP2004242513-A.
PD 02-SEP-2004.
PA (DOKU-) DOKURITSU GYOSEI HOJIN KAGAKU GIJUTSU SH.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 480
ID ADR46571 standard; protein; 299 AA.
DE Human JAM-1, F11 receptor (F11R) transcript variant 4.
PN JP2004242513-A.
PD 02-SEP-2004.
PA (DOKU-) DOKURITSU GYOSEI HOJIN KAGAKU GIJUTSU SH.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 481
ID ADR46573 standard; protein; 299 AA.
DE Human JAM-1, F11 receptor (F11R) transcript variant 4.
PN JP2004242513-A.
PD 02-SEP-2004.
PA (DOKU-) DOKURITSU GYOSEI HOJIN KAGAKU GIJUTSU SH.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 482
ID ADR46579 standard; protein; 299 AA.
DE Human JAM-1, F11 receptor (F11R) transcript variant 5.
PN JP2004242513-A.
PD 02-SEP-2004.
PA (DOKU-) DOKURITSU GYOSEI HOJIN KAGAKU GIJUTSU SH.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 483
ID ADT03591 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003152922-A1.
PD 14-AUG-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 484
ID ADT94260 standard; protein; 299 AA.
DE Human PRO301 protein.
PN AU2003259607-A1.
PD 27-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;

RESULT 485
ID ADS74554 standard; protein; 299 AA.
DE Human secreted/transmembrane protein #25.
PN US2004185531-A1.
PD 23-SEP-2004.
PA (ASHK//) ASHKENAZI A.
PA (BOTS//) BOTSTEIN D.
PA (DESN//) DESNOYERS L.
PA (EATO//) EATON D L.
PA (FERR//) FERRARA N.
PA (FILV//) FILVAROFF E.
PA (FONG//) FONG S.
PA (GAOW//) GAO W.
PA (GERB//) GERBER H.
PA (GERR//) GERRITSEN M E.
PA (GODD//) GODDARD A.
PA (GODO//) GODOWSKI P J.
PA (GRIM//) GRIMALDI C J.
PA (GURN//) GURNEY A L.
PA (HILL//) HILLAN K J.
PA (KLJA//) KLJAVIN I J.
PA (MATH//) MATHER J P.
PA (PANJ//) PAN J.
PA (PAON//) PAONI N F.
PA (ROYM//) ROY M A.
PA (STEW//) STEWART T A.
PA (TUMA//) TUMAS D.
PA (WILL//) WILLIAMS P M.
PA (WOOD//) WOOD W I.
Query Match 10.6%; Score 178.5; DB 8; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.1e-06;
RESULT 486
ID ADE09073 standard; protein; 320 AA.
DE Novel protein-related contig polypeptide sequence #139.
PN WO2003054152-A2.
PD 03-JUL-2003.
PA (HYSE-) HYSEQ INC.
Query Match 10.6%; Score 178.5; DB 7; Length 320;
Best Local Similarity 26.5%; Pred. No. 1.2e-06;
RESULT 487
ID ADU67616 standard; protein; 335 AA.
DE Human ovarian specific polypeptide SEQ ID NO:330.
PN WO2004013311-A2.
PD 12-FEB-2004.
PA (DIAD-) DIADEXUS INC.
Query Match 10.6%; Score 178.5; DB 8; Length 335;
Best Local Similarity 26.5%; Pred. No. 1.3e-06;
RESULT 488
ID ADR08038 standard; protein; 336 AA.
DE Novel protein (useful for identifying genetic disorders) #193.
PN WO2003054152-A2.
PD 03-JUL-2003.
PA (HYSE-) HYSEQ INC.
Query Match 10.6%; Score 178.5; DB 7; Length 336;
Best Local Similarity 26.5%; Pred. No. 1.3e-06;
RESULT 489
ID AAW14146 standard; protein; 319 AA.
DE Human A33 antigen.
PN WO9708189-A1.
PD 06-MAR-1997.
PA (LUDW-) LUDWIG INST CANCER RES.
Query Match 10.5%; Score 177; DB 2; Length 319;
Best Local Similarity 27.5%; Pred. No. 1.6e-06;
RESULT 490
ID AAY23323 standard; protein; 319 AA.
DE Amino acid sequence of the A33 antigen.
PN WO9927098-A2.
PD 03-JUN-1999.
PA (GETH) GENENTECH INC.
Query Match 10.5%; Score 177; DB 2; Length 319;
Best Local Similarity 27.5%; Pred. No. 1.6e-06;
RESULT 491
ID AAB65863 standard; protein; 319 AA.


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DE Human A33 protein SEQ ID NO: 67.
PN WO200078808-A1.
PA (MILL-) MILLENNIUM PHARM INC.
Query Match 10.5%; Score 177; DB 4; Length 319;
Best Local Similarity 27.5%; Pred. No. 1.6e-06;
RESULT 492
ID ADA10947 standard; protein; 319 AA.
DE Human cDNA differentially expressed in colon cancer #43 product.
PN US2002160382-A1.
PD 31-OCT-2002.
PA (LASE/) LASEK A W.
PA (JONE/) JONES D A.
Query Match 10.5%; Score 177; DB 6; Length 319;
Best Local Similarity 27.5%; Pred. No. 1.6e-06;
RESULT 493
ID ADH62533 standard; protein; 319 AA.
DE Human A33 antigenic protein.
PN US2003171568-A1.
PD 11-SEP-2003.
PA (ASHK/) ASHKENAZI A.
PA (FONG/) FONG S.
PA (GODD/) GODDARD A.
PA (GURN/) GURNEY A L.
PA (NAPI/) NAPIER M A.
PA (TUMA/) TUMAS D.
PA (WOOD/) WOOD W I.
Query Match 10.5%; Score 177; DB 7; Length 319;
Best Local Similarity 27.5%; Pred. No. 1.6e-06;
RESULT 494
ID ADN39847 standard; protein; 319 AA.
DE Cancer/angiogenesis/fibrosis-related polypeptide, SEQ ID NO:C217.
PN WO2003042661-A2.
PD 22-MAY-2003.
PA (EOSB-) EOS BIOTECHNOLOGY INC.
Query Match 10.5%; Score 177; DB 7; Length 319;
Best Local Similarity 27.5%; Pred. No. 1.6e-06;
RESULT 495
ID ADN35289 standard; protein; 319 AA.
DE Human A33 antigen protein.
PN WO2004031105-A2.
PD 15-APR-2004.
PA (GETH ) GENENTECH INC.
Query Match 10.5%; Score 177; DB 8; Length 319;
Best Local Similarity 27.5%; Pred. No. 1.6e-06;
RESULT 496
ID ADP54587 standard; protein; 319 AA.
DE Human PRO protein sequence SEQ ID NO:563.
PN WO2004039956-A2.
PD 13-MAY-2004.
PA (GETH ) GENENTECH INC.
Query Match 10.5%; Score 177; DB 8; Length 319;
Best Local Similarity 27.5%; Pred. No. 1.6e-06;
RESULT 497
ID ABP62881 standard; protein; 336 AA.
DE Human polypeptide SEQ ID NO 318.
PN WO200218424-A2.
PD 07-MAR-2002.
PA (HYSE-) HYSEQ INC.
Query Match 10.5%; Score 177; DB 5; Length 336;
Best Local Similarity 27.5%; Pred. No. 1.7e-06;
RESULT 498
ID ADC78439 standard; protein; 299 AA.
DE Human PRO301 protein.
PN WO200015796-A2.
PD 23-MAR-2000.
PA (GETH ) GENENTECH INC.
Query Match 10.5%; Score 176.5; DB 3; Length 299;
Best Local Similarity 26.5%; Pred. No. 1.6e-06;
RESULT 499
ID AAY08073 standard; protein; 268 AA.
DE Human A33 protein.
PN WO9914241-A2.

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PD 25-MAR-1999.
PA (GETH ) GENENTECH INC.
Query Match 10.4%; Score 176; DB 2; Length 268;
Best Local Similarity 27.7%; Pred. No. 1.6e-06;
RESULT 500
ID ADH62551 standard; protein; 268 AA.
DE Human A33 antigenic protein fragment #1.
PN US2003171568-A1.
PD 11-SEP-2003.
PA (ASHK/) ASHKENAZI A.
PA (FONG/) FONG S.
PA (GODD/) GODDARD A.
PA (GURN/) GURNEY A L.
PA (NAPI/) NAPIER M A.
PA (TUMA/) TUMAS D.
PA (WOOD/) WOOD W I.
Query Match 10.4%; Score 176; DB 7; Length 268;
Best Local Similarity 27.7%; Pred. No. 1.6e-06;
RESULT 501
ID AAY23327 standard; protein; 270 AA.
DE An A33 related antigen sequence.
PN WO9927098-A2.
PD 03-JUN-1999.
PA (GETH ) GENENTECH INC.
Query Match 10.4%; Score 176; DB 2; Length 270;
Best Local Similarity 27.7%; Pred. No. 1.6e-06;
RESULT 502
ID AAY23329 standard; protein; 273 AA.
DE An A33 related antigen sequence.
PN WO9927098-A2.
PD 03-JUN-1999.
PA (GETH ) GENENTECH INC.
Query Match 10.4%; Score 176; DB 2; Length 273;
Best Local Similarity 27.7%; Pred. No. 1.6e-06;
RESULT 503
ID AAY08075 standard; protein; 273 AA.
DE Human A33 protein fragment #2.
PN WO9914241-A2.
PD 25-MAR-1999.
PA (GETH ) GENENTECH INC.
Query Match 10.4%; Score 176; DB 2; Length 273;
Best Local Similarity 27.7%; Pred. No. 1.6e-06;
RESULT 504
ID ADH62553 standard; protein; 273 AA.
DE Human A33 antigenic protein fragment #2.
PN US2003171568-A1.
PD 11-SEP-2003.
PA (ASHK/) ASHKENAZI A.
PA (FONG/) FONG S.
PA (GODD/) GODDARD A.
PA (GURN/) GURNEY A L.
PA (NAPI/) NAPIER M A.
PA (TUMA/) TUMAS D.
PA (WOOD/) WOOD W I.
Query Match 10.4%; Score 176; DB 7; Length 273;
Best Local Similarity 27.7%; Pred. No. 1.6e-06;
RESULT 505
ID ADH80722 standard; protein; 301 AA.
DE Human polypeptide #39.
PN US2003232054-A1.
PD 18-DEC-2003.
PA (TANG/) TANG Y T.
PA (LIUC/) LIU C.
PA (ASUN/) ASUNDI V.
PA (CHEN/) CHEN R.
PA (QIAN/) QIAN X B.
PA (WEHR/) WEHRMAN T.
PA (ZHAN/) ZHANG J.
PA (ZHOU/) ZHOU P.
PA (CAOY/) CAO Y.
PA (DRMA/) DRMANAC R T.
Query Match 10.3%; Score 173.5; DB 8; Length 301;

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Best Local Similarity 26.6%; Pred. No. 3e-06;
RESULT 506
ID ADK40854 standard; protein; 316 AA.
DE Human A33 molecule.
PN US6699688-B1.
PD 02-MAR-2004.
PA (UUNY) UNIV NEW YORK STATE RES FOUND.
Query Match 10.2%; Score 172.5; DB 8; Length 316;
Best Local Similarity 26.4%; Pred. No. 3.9e-06;
RESULT 507
ID ADQ89964 standard; protein; 34350 AA.
DE Antagonist of cell cycle progression polypeptide #197.
PN WO2004063362-A2.
PD 29-JUL-2004.
PA (CYCL-) CYCLACEL LTD.
Query Match 10.2%; Score 172; DB 8; Length 34350;
Best Local Similarity 27.5%; Pred. No. 0.0024;
RESULT 508
ID AAY23326 standard; protein; 260 AA.
DE An A33 related antigen sequence.
PN WO927098-A2.
PD 03-JUN-1999.
PA (GETH) GENENTECH INC.
Query Match 10.0%; Score 169.5; DB 2; Length 260;
Best Local Similarity 26.1%; Pred. No. 5.4e-06;
RESULT 509
ID AAY08072 standard; protein; 260 AA.
DE Human DNA40628 protein.
PN WO9914241-A2.
PD 25-MAR-1999.
PA (GETH) GENENTECH INC.
Query Match 10.0%; Score 169.5; DB 2; Length 260;
Best Local Similarity 26.1%; Pred. No. 5.4e-06;
RESULT 510
ID ADHG2550 standard; protein; 260 AA.
DE Human PRO301 protein fragment #1.
PN US2003171568-A1.
PD 11-SEP-2003.
PA (ASHK/) ASHKENAZI A.
PA (FONG/) FONG S.
PA (GODD/) GODDARD A.
PA (GURN/) GURNEY A L.
PA (NAPI/) NAPIER M A.
PA (TUMA/) TUMAS D.
PA (WOOD/) WOOD W I.
Query Match 10.0%; Score 169.5; DB 7; Length 260;
Best Local Similarity 26.1%; Pred. No. 5.4e-06;
RESULT 511
ID AAY23328 standard; protein; 263 AA.
DE An A33 related antigen sequence.
PN WO927098-A2.
PD 03-JUN-1999.
PA (GETH) GENENTECH INC.
Query Match 10.0%; Score 169.5; DB 2; Length 263;
Best Local Similarity 26.1%; Pred. No. 5.5e-06;
RESULT 512
ID AAY08074 standard; protein; 263 AA.
DE Human DNA40628 protein fragment #2.
PN WO9914241-A2.
PD 25-MAR-1999.
PA (GETH) GENENTECH INC.
Query Match 10.0%; Score 169.5; DB 2; Length 263;
Best Local Similarity 26.1%; Pred. No. 5.5e-06;
RESULT 513
ID ADHG2552 standard; protein; 263 AA.
DE Human PRO301 protein fragment #2.
PN US2003171568-A1.
PD 11-SEP-2003.
PA (ASHK/) ASHKENAZI A.
PA (FONG/) FONG S.
PA (GODD/) GODDARD A.
PA (GURN/) GURNEY A L.
PA (NAPI/) NAPIER M A.
PA (TUMA/) TUMAS D.
PA (WOOD/) WOOD W I.
Query Match 10.0%; Score 169.5; DB 7; Length 263;
Best Local Similarity 26.1%; Pred. No. 5.4e-06;
RESULT 514
ID ADR46581 standard; protein; 300 AA.
DE Mouse junctional adhesion molecule-1, SEQ ID 12.
PN JP2004242513-A.
PD 02-SEP-2004.
PA (DOKU-) DOKURITSU GYOSEI HOJIN KAKAKU GIJUTSU SH.
Query Match 10.0%; Score 168; DB 8; Length 300;
Best Local Similarity 24.8%; Pred. No. 8.8e-06;
RESULT 515
ID AAW61379 standard; protein; 298 AA.
DE Human junctional adhesion molecule protein.
PN WO9824897-A1.
PD 11-JUN-1998.
PA (HOFF) HOFFMANN LA ROCHE & CO AG F.
Query Match 9.9%; Score 167; DB 2; Length 298;
Best Local Similarity 24.8%; Pred. No. 1.1e-05;
RESULT 516
ID AAW61380 standard; protein; 300 AA.
DE Mouse junctional adhesion molecule protein.
PN WO9824897-A1.
PD 11-JUN-1998.
PA (HOFF) HOFFMANN LA ROCHE & CO AG F.
Query Match 9.9%; Score 167; DB 2; Length 300;
Best Local Similarity 24.8%; Pred. No. 1.1e-05;
RESULT 517
ID AAY23325 standard; protein; 300 AA.
DE A33 related antigen JAM.
PN WO927098-A2.
PD 03-JUN-1999.
PA (GETH) GENENTECH INC.
Query Match 9.9%; Score 167; DB 2; Length 300;
Best Local Similarity 24.8%; Pred. No. 1.1e-05;
RESULT 518
ID ADHG2537 standard; protein; 300 AA.
DE Murine JAM protein used in the exemplification of the invention.
PN US2003171568-A1.
PD 11-SEP-2003.
PA (ASHK/) ASHKENAZI A.
PA (FONG/) FONG S.
PA (GODD/) GODDARD A.
PA (GURN/) GURNEY A L.
PA (NAPI/) NAPIER M A.
PA (TUMA/) TUMAS D.
PA (WOOD/) WOOD W I.
Query Match 9.9%; Score 167; DB 7; Length 300;
Best Local Similarity 24.8%; Pred. No. 1.1e-05;
RESULT 519
ID ADK40853 standard; protein; 300 AA.
DE Mouse junction adhesion molecule (JAM).
PN US6699688-B1.
PD 02-MAR-2004.
PA (UUNY) UNIV NEW YORK STATE RES FOUND.
Query Match 9.9%; Score 167; DB 8; Length 300;
Best Local Similarity 24.8%; Pred. No. 1.1e-05;
RESULT 520
ID ADN35293 standard; protein; 300 AA.
DE Human JAM protein.
PN WO2004031105-A2.
PD 15-APR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.9%; Score 167; DB 8; Length 300;
Best Local Similarity 24.8%; Pred. No. 1.1e-05;
RESULT 521
ID ABB83928 standard; protein; 365 AA.
DE DCAR SEQ ID NO 4.
PN US2002059654-A1.
PD 16-MAY-2002.
PA (BUHL/) BUHLER T.
PA (GADI/) GADIENT R A.

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PA (KORN/) KORN R.
PA (MOVV/) MOVVA R.
Query Match
Best Local Similarity 9.8%; Score 166; DB 5; Length 365;
Best Local Similarity 24.2%; Pred. No. 1.7e-05;
RESULT 522
ID ABB83927 standard; protein; 261 AA.
DE C-terminally truncated pCAR SEQ ID NO 2.
PN US2002059654-A1.
PD 16-MAY-2002.
PA (BUHL/) BUHLER T.
PA (GADI/) GADIENT R. A.
PA (KORN/) KORN R.
PA (MOVV/) MOVVA R.
Query Match
Best Local Similarity 9.7%; Score 163.5; DB 5; Length 261;
Best Local Similarity 27.7%; Pred. No. 1.8e-05;
RESULT 523
ID AAM14158 standard; protein; 318 AA.
DE Mouse A33 antigen.
PN WO9708189-A1.
PD 06-MAR-1997.
PA (LUDW-) LUDWIG INST CANCER RES.
Query Match
Best Local Similarity 9.6%; Score 162.5; DB 2; Length 318;
Best Local Similarity 22.9%; Pred. No. 2.8e-05;
RESULT 524
ID AAB39253 standard; protein; 280 AA.
DE Gene 15 human secreted protein homologous amino acid sequence #133.
PN WO200056754-A1.
PD 28-SEP-2000.
PA (HUNA-) HUMAN GENOME SCI INC.
Query Match
Best Local Similarity 9.5%; Score 160; DB 3; Length 280;
Best Local Similarity 24.7%; Pred. No. 3.9e-05;
RESULT 525
ID ADA50171 standard; protein; 412 AA.
DE Human CAR/mouse SCF fusion protein.
PN US2003092068-A1.
PD 15-MAY-2003.
PA (ITO/) ITOH A.
PA (HANA/) HANAZONO Y.
PA (OKAD/) OKADA T.
PA (OZAW/) OZAWA K.
Query Match
Best Local Similarity 9.4%; Score 158.5; DB 6; Length 412;
Best Local Similarity 24.8%; Pred. No. 8.7e-05;
RESULT 526
ID AAY72878 standard; protein; 352 AA.
DE Human PRO5723 protein encoded by DNA82361 cDNA clone.
PN WO200116319-A2.
PD 08-MAR-2001.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.4%; Score 158; DB 4; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 527
ID AAB50930 standard; protein; 352 AA.
DE Human PRO5723 protein.
PN WO200073452-A2.
PD 07-DEC-2000.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.4%; Score 158; DB 4; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 528
ID AAB65294 standard; protein; 352 AA.
DE Human PRO5723 protein sequence SEQ ID NO:505.
PN WO200073454-A1.
PD 07-DEC-2000.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.4%; Score 158; DB 4; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 529
ID ABB84956 standard; protein; 352 AA.
DE Human PRO5723 protein sequence SEQ ID NO:280.
PN WO200200690-A2.
PD 03-JAN-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.4%; Score 158; DB 5; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 530
ID ABB95562 standard; protein; 352 AA.
DE Human angiogenesis related protein PRO5723 SEQ ID NO: 280.
PN WO200208284-A2.
PD 31-JAN-2002.
PA (GETH) GENENTECH INC.
PA (BAKE/) BAKER K P.
PA (FERR/) FERRARA N.
PA (GERB/) GERBER H.
PA (GERR/) GERRITSEN M E.
PA (GODD/) GODDARD A.
PA (GODO/) GODOWSKI P J.
PA (GURN/) GURNEY A L.
PA (HILL/) HILLAN K J.
PA (MARS/) MARSTERS S A.
PA (PANJ/) PAN J.
PA (PAON/) PAONI N P.
PA (STEP/) STEPHAN J F.
PA (WATA/) WATANABE C K.
PA (WILL/) WILLIAMS P M.
PA (WOOD/) WOOD W I.
Query Match
Best Local Similarity 9.4%; Score 158; DB 5; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 531
ID ABU58109 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003027163-A1.
PD 06-FEB-2003.
Query Match
Best Local Similarity 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 532
ID ABU59187 standard; protein; 352 AA.
DE Novel human secreted or transmembrane protein PRO5723.
PN US2002132252-A1.
PD 19-SEP-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 533
ID ABU82699 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003032023-A1.
PD 13-FEB-2003.
Query Match
Best Local Similarity 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 534
ID ABU60618 standard; protein; 352 AA.
DE Human secreted/transmembrane protein, #177.
PN US2002160384-A1.
PD 31-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 535
ID ABU80846 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003036635-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 536
ID ABO33812 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003045687-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 537
ID ABU14000 standard; protein; 352 AA.
DE Human PRO5723 polypeptide.
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PN US2002103125-A1.
PD 01-AUG-2002.
PA (GETH) GENENTECH LTD. 9.4%; Score 158; DB 6; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 538
ID ABU72595 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US200303531-A1.
PD 02-JAN-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 539
ID ABG74762 standard; protein; 352 AA.
DE Human PRO5723 protein.
PN US2002192752-A1.
PD 19-DEC-2002.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 540
ID ABU59334 standard; protein; 352 AA.
DE Human secreted/transmembrane protein, #177.
PN US2003027162-A1.
PD 06-FEB-2003.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 541
ID ABO26031 standard; protein; 352 AA.
DE Human PRO5723 polypeptide.
PN US2002127576-A1.
PD 12-SEP-2002.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 542
ID ABU82155 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US200308063-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 543
ID ABU59040 standard; protein; 352 AA.
DE Human secreted/transmembrane protein, #177.
PN US2002142961-A1.
PD 03-OCT-2002.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 544
ID ABU92418 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003022187-A1.
PD 30-JAN-2003.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 545
ID ABU59483 standard; protein; 352 AA.
DE Novel human secreted or transmembrane protein PRO3301.
PN US2003027985-A1.
PD 06-FEB-2003.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 546
ID ABU92249 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003017476-A1.
PD 23-JAN-2003.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;

RESULT 547
ID ABU10955 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2002123463-A1.
PD 05-SEP-2002.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 548
ID ABU81707 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2002177164-A1.
PD 28-NOV-2002.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 549
ID ABU8646 standard; protein; 352 AA.
DE Human secreted and transmembrane polypeptide PRO5723.
PN US2002197615-A1.
PD 26-DEC-2002.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 550
ID ABO34160 standard; protein; 352 AA.
DE Human PRO5723 polypeptide.
PN US2003017981-A1.
PD 23-JAN-2003.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 551
ID ABJ72335 standard; protein; 352 AA.
DE Human PRO5723 protein.
PN US2003050448-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 552
ID ADA38016 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003008297-A1.
PD 09-JAN-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 6; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 553
ID ADA21702 standard; protein; 352 AA.
DE Human secreted/transmembrane polypeptide PRO5723.
PN US2003054404-A1.
PD 20-MAR-2003.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 554
ID ADA10489 standard; protein; 352 AA.
DE Human secreted/transmembrane protein, PRO5723.
PN US2003059831-A1.
PD 27-MAR-2003.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 555
ID ADA18033 standard; protein; 352 AA.
DE Human PRO5723 polypeptide.
PN US2003054987-A1.
PD 20-MAR-2003.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 556
ID ADA28141 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003054359-A1.
PD 20-MAR-2003.

Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 557
ID ADA94721 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003059832-A1.
PD 27-MAR-2003.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 558
ID ADA38946 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003059780-A1.
PD 27-MAR-2003.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 559
ID ABJ72463 standard; protein; 352 AA.
DE Human PRO5723 protein.
PN US2003027988-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 560
ID ADA93067 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003060407-A1.
PD 27-MAR-2003.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 561
ID ABO34358 standard; protein; 352 AA.
DE Human secreted/transmembrane polypeptide PRO 5723.
PN US2003044934-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 6; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 562
ID ABO53246 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003044806-A1.
PD 06-MAR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 563
ID ADA22628 standard; protein; 352 AA.
DE Human secreted/transmembrane polypeptide PRO5723.
PN US2003040473-A1.
PD 27-FEB-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 564
ID ABO22616 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003017982-A1.
PD 23-JAN-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 565
ID ADA06794 standard; protein; 352 AA.
DE Human secreted/transmembrane PRO polypeptide #141.
PN US2003049638-A1.
PD 13-MAR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 566
ID ABJ72165 standard; protein; 352 AA.
DE Human membrane bound receptor/protein PRO5723 amino acid sequence.
PN US2003065147-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.

Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 567
ID ADA39487 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003059782-A1.
PD 27-MAR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 568
ID ADB83706 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003073814-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 569
ID ADB80812 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US200308068-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 570
ID ADB73353 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003096968-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 571
ID ADB96513 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003054403-A1.
PD 20-MAR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 572
ID ADB78435 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003092889-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 573
ID ADB85083 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003073817-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 574
ID ADB78189 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003092886-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 575
ID ADB87255 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US200308067-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 576
ID ADB72165 standard; protein; 352 AA.
DE Human membrane bound receptor/protein PRO5723 amino acid sequence.
PN US2003065147-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.

ID ADB84837 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003092890-A1.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 577
ID ADB83952 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003069397-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 578
ID ADB73107 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003092887-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 579
ID ADC57985 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003027754-A1.
PD 06-FEB-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 580
ID ADC55349 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003045463-A1.
PD 06-MAR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 581
ID ADC12216 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003049681-A1.
PD 13-MAR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 582
ID ADC56638 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003064375-A1.
PD 03-APR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 583
ID ADC11683 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003069403-A1.
PD 10-APR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 584
ID ADC36945 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003088065-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 585
ID ADC21935 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003096969-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 586
ID ADC49966 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003088064-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 587
ID ADC49165 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003088070-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 588
ID ADC49682 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003088071-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 589
ID ADC47543 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003088072-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 590
ID ADC14805 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003082546-A1.
PD 01-MAY-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 591
ID ADC47288 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003105288-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 592
ID ADD08337 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003068623-A1.
PD 10-APR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 593
ID ADC82162 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003083461-A1.
PD 01-MAY-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 594
ID ADD07804 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2002193299-A1.
PD 19-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 595
ID ADC78163 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.

PN US2003096972-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 596
ID ADC82695 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003059833-A1.
PD 27-MAR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 597
ID ADD06398 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003073816-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 598
ID ADD10569 standard; protein; 352 AA.
DE Human secreted/transmembrane PRO polypeptide #140.
PN US2003105011-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 599
ID ADD08875 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003073090-A1.
PD 17-APR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 600
ID ADC77917 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US200308066-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 601
ID ADD07124 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2002193300-A1.
PD 19-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 602
ID ADD11529 standard; protein; 352 AA.
DE Human secreted/transmembrane PRO polypeptide #140.
PN US2003105013-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 603
ID ADC83371 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003059783-A1.
PD 27-MAR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 604
ID ADD50880 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003105291-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 605
ID ADD15380 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003059437-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 606
ID ADD51126 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003105290-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 607
ID ADD55478 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003077593-A1.
PD 24-APR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 608
ID ADD37322 standard; protein; 352 AA.
DE Human secreted/transmembrane PRO polypeptide #140.
PN US2003105012-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 609
ID ADD56436 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003077594-A1.
PD 24-APR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 610
ID ADD50607 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003096971-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 611
ID ADD54874 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2002132253-A1.
PD 19-SEP-2002.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 612
ID ADD50361 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003096970-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 613
ID ADD51372 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003105289-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 614
ID ADE31893 standard; protein; 352 AA.

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DE Human secreted/transmembrane protein PRO5723.
PN US2003068647-A1.
PD 10-APR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 615
ID ADE27028 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003087304-A1.
PD 08-MAY-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 616
ID ADE26495 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003087305-A1.
PD 08-MAY-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 617
ID ADE67432 standard; protein; 352 AA.
DE Human PRO5723 amino acid sequence SEQ ID NO:505.
PN US2002198148-A1.
PD 26-DEC-2002.
PA (GETH ) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 618
ID ADE94085 standard; protein; 352 AA.
DE Immune disease treatment/diagnosis related PRO5723.
PN US2003082199-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 619
ID ADI35686 standard; protein; 352 AA.
DE Human PRO polypeptide #141.
PN US2003050457-A1.
PD 13-MAR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 620
ID ADI00179 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003049682-A1.
PD 13-MAR-2003.
Query Match 9.4%; Score 158; DB 7; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 621
ID ADC48919 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003092888-A1.
PD 15-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 622
ID ADE21090 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100735-A1.
PD 29-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 623
ID ADE05934 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100728-A1.
PD 29-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 624
ID ADD75163 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100712-A1.
PD 29-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 625
ID ADD75909 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100717-A1.
PD 29-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 626
ID ADD85141 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100722-A1.
PD 29-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 627
ID ADD86967 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100738-A1.
PD 29-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 628
ID ADE20844 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100734-A1.
PD 29-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 629
ID ADE39141 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003096362-A1.
PD 22-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 630
ID ADE05688 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100727-A1.
PD 29-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 631
ID ADD73673 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100711-A1.
PD 29-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 632
ID ADD78513 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100737-A1.
PD 29-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 633
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ID ADE41530 standard; protein; 352 AA.
DE Human secreted/transmembrane PRO polypeptide #140.
PN US2003100497-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 634
ID ADE21336 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100736-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 635
ID ADD77451 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100732-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 636
ID ADE20598 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100733-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 637
ID ADD75663 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100064-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 638
ID ADD74179 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100708-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 639
ID ADD74425 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100709-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 640
ID ADD76155 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100718-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 641
ID ADD85647 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100721-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 642
ID ADE05196 standard; protein; 352 AA.

DE Human PRO polypeptide #108.
PN US2003100726-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 643
ID ADD75409 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100714-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 644
ID ADD76953 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100715-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 645
ID ADD86721 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100719-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 646
ID ADD78189 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100731-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 647
ID ADD7697 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100729-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 648
ID ADD77943 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100730-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 649
ID ADD85401 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100725-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 650
ID ADD73933 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100710-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 651
ID ADD74671 standard; protein; 352 AA.
DE Human PRO polypeptide #108.

PN US2003100713-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 652
ID ADG77199 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100716-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 653
ID ADG85893 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003100720-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 654
ID ADE05442 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100723-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 655
ID ADD74917 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003100724-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 656
ID ADF35631 standard; protein; 352 AA.
DE Human PRO5723 polypeptide.
PN US2003194760-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 657
ID ADG11881 standard; protein; 352 AA.
DE Human PRO5723 polypeptide.
PN US2003228655-A1.
PD 11-DEC-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 658
ID ADG05729 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003096959-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 659
ID ADG27283 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003096962-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 660
ID ADG11346 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003096967-A1.
PD 22-MAY-2003.

PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 661
ID ADG12125 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003096963-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 662
ID ADF94682 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003096964-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 663
ID ADG06778 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003096966-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 664
ID ADH39122 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003096965-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 665
ID ADH19751 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003228656-A1.
PD 11-DEC-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 666
ID ADH21244 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003224358-A1.
PD 04-DEC-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 667
ID ADH20284 standard; protein; 352 AA.
DE Human secreted/transmembrane protein PRO5723.
PN US2003219856-A1.
PD 27-NOV-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 668
ID ADH43713 standard; protein; 352 AA.
DE Human PRO polypeptide #140.
PN US2003224984-A1.
PD 04-DEC-2003.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 669
ID ADG34212 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2004006206-A1.
PD 08-JAN-2004.
PA (GETH) GENENTECH INC. 9.4%; Score 158; DB 8; Length 352;
Query Match 23.3%; Pred. No. 7.8e-05;

Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 670
ID ADI33682 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2003096960-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 671
ID ADH69776 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US2004019183-A1.
PD 29-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 672
ID ADI29937 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US2003096961-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 673
ID ADM27334 standard; protein; 352 AA.
DE Novel human secreted and transmembrane protein PRO5723.
PN US200404179-A1.
PD 04-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 674
ID ADR83058 standard; protein; 352 AA.
DE Human PRO polypeptide #140.
PN US2004043927-A1.
PD 04-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 675
ID ADK66692 standard; protein; 352 AA.
DE Human PRO polypeptide #108.
PN US200404180-A1.
PD 04-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.4%; Score 158; DB 8; Length 352;
Best Local Similarity 23.3%; Pred. No. 7.8e-05;
RESULT 676
ID AAW69697 standard; protein; 365 AA.
DE Human coxsackievirus and Ad2 and Ad5 receptor HCAR protein.
PN WO9833819-A1.
PD 06-AUG-1998.
PA (UUNY) UNIV NEW YORK STATE.
Query Match 9.4%; Score 158; DB 2; Length 365;
Best Local Similarity 23.3%; Pred. No. 8.2e-05;
RESULT 677
ID AAW57212 standard; protein; 365 AA.
DE Human coxsackievirus and adenovirus receptor.
PN WO9811221-A2.
PD 19-MAR-1998.
PA (DAND) DANA FARBER CANCER INST INC.
Query Match 9.4%; Score 158; DB 2; Length 365;
Best Local Similarity 23.3%; Pred. No. 8.2e-05;
RESULT 678
ID AAB47270 standard; protein; 365 AA.
DE Human CAR.
PN US6245966-B1.
PD 12-JUN-2001.
PA (UYTE-) UNIV TECHNOLOGY CORP.
Query Match 9.4%; Score 158; DB 4; Length 365;
Best Local Similarity 23.3%; Pred. No. 8.2e-05;

RESULT 679
ID ABB08040 standard; protein; 365 AA.
DE Human coxsackie-adenovirus receptor (CAR).
PN WO200229072-A2.
PD 11-APR-2002.
PA (NOVS) NOVARTIS AG.
PA (NOVS) NOVARTIS-ERFINDUNGEN VERW GES MBH.
Query Match 9.4%; Score 158; DB 5; Length 365;
Best Local Similarity 23.3%; Pred. No. 8.2e-05;
RESULT 680
ID ABJ37063 standard; protein; 365 AA.
DE Human breast cancer / ovarian cancer related protein #39.
PN WO2003000012-A2.
PD 03-JAN-2003.
PA (MILL-) MILLENNIUM PHARM INC.
Query Match 9.4%; Score 158; DB 6; Length 365;
Best Local Similarity 23.3%; Pred. No. 8.2e-05;
RESULT 681
ID AD897544 standard; protein; 365 AA.
DE Human CAR wild-type protein.
PN WO2003070915-A2.
PD 28-AUG-2003.
PA (USSH) US DEPT HEALTH & HUMAN SERVICES.
Query Match 9.4%; Score 158; DB 7; Length 365;
Best Local Similarity 23.3%; Pred. No. 8.2e-05;
RESULT 682
ID ADN95226 standard; protein; 365 AA.
DE Human BEC/LEC-related protein sequence SeqID148.
PN WO2003080640-A1.
PD 02-OCT-2003.
PA (LUDW-) LUDWIG INST CANCER RES.
PA (LICN) LICENTIA LTD.
Query Match 9.4%; Score 158; DB 7; Length 365;
Best Local Similarity 23.3%; Pred. No. 8.2e-05;
RESULT 683
ID ABUL2046 standard; protein; 505 AA.
DE Human NOV4a CG59871-01 protein SEQ ID 12.
PN WO200281625-A2.
PD 17-OCT-2002.
PA (CURA-) CURAGEN CORP.
Query Match 9.4%; Score 158; DB 6; Length 505;
Best Local Similarity 23.3%; Pred. No. 0.00013;
RESULT 684
ID ADJ67617 standard; protein; 351 AA.
DE Human ovarian specific polypeptide SEQ ID NO:331.
PN WO200401311-A2.
PD 12-FEB-2004.
PA (DIAD-) DIADEXUS INC.
Query Match 9.3%; Score 157.5; DB 8; Length 351;
Best Local Similarity 23.0%; Pred. No. 8.6e-05;
RESULT 685
ID ADA50172 standard; protein; 493 AA.
DE Human CAR/mouse anti-CD34 antibody fusion protein.
PN US2003092068-A1.
PD 15-MAY-2003.
PA (ITOH/) ITOH A.
PA (HANA/) HANAZONO Y.
PA (OKAD/) OKADA T.
PA (OZAW/) OZAWA K.
Query Match 9.3%; Score 157.5; DB 6; Length 493;
Best Local Similarity 23.7%; Pred. No. 0.00014;
RESULT 686
ID AAW82729 standard; protein; 264 AA.
DE Adenovirus pACTSG2-SCAR protein.
PN WO9854346-A1.
PD 03-DEC-1998.
PA (GENV-) GENVEC INC.
Query Match 9.3%; Score 156.5; DB 2; Length 264;
Best Local Similarity 25.9%; Pred. No. 7.1e-05;
RESULT 687
ID AAW82730 standard; protein; 277 AA.
DE Adenovirus SCAR.RGD protein.
PN WO9854346-A1.

PD 03-DEC-1998.
PA (GENV-) GENVEC INC. 9.3%; Score 156.5; DB 2; Length 277;
Best Local Similarity 25.9%; Pred. No. 7.6e-05;
RESULT 688
ID AAU83699 standard; protein; 290 AA.
DE Human PRO protein, Seq ID No 216.
PN WO200208288-A2.
PD 31-JAN-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 25.9%; Score 156.5; DB 5; Length 290;
RESULT 689
ID AAU82731 standard; protein; 397 AA.
DE Adenovirus PACSG2SCAR.sig chimeric protein.
PN WO9854346-A1.
PD 03-DEC-1998.
PA (GENV-) GENVEC INC.
Query Match
Best Local Similarity 25.9%; Score 156.5; DB 2; Length 397;
RESULT 690
ID ADA50170 standard; protein; 412 AA.
DE Human CAR/SCF fusion protein.
PN US2003092068-A1.
PD 15-MAY-2003.
PA (ITOH/) ITOH A.
PA (HANA/) HANAZONO Y.
PA (OKAD/) OKADA T.
PA (OZAW/) OZAWA K.
Query Match
Best Local Similarity 25.9%; Score 156.5; DB 6; Length 412;
RESULT 691
ID ABG02019 standard; protein; 737 AA.
DE Novel human diagnostic protein #2010.
PN WO200175067-A2.
PD 11-OCT-2001.
PA (HYSE-) HYSEQ INC.
Query Match
Best Local Similarity 24.3%; Score 156; DB 4; Length 737;
RESULT 692
ID ADK66297 standard; protein; 358 AA.
DE Human prostatic carcinoma derived protein SEQ ID 151 #2.
PN WO2004076614-A2.
PD 10-SEP-2004.
PA (HINZ/) HINZMANN B.
PA (DAHL/) DAHL E.
PA (ROSE/) ROSENTHAL A.
PA (HERM/) HERMANN K.
PA (PILA/) PILARSKY C.
Query Match
Best Local Similarity 23.1%; Score 153.5; DB 8; Length 358;
RESULT 693
ID ADK66858 standard; protein; 358 AA.
DE Human prostatic carcinoma derived DNA SEQ ID 151 #4.
PN WO2004076614-A2.
PD 10-SEP-2004.
PA (HINZ/) HINZMANN B.
PA (DAHL/) DAHL E.
PA (ROSE/) ROSENTHAL A.
PA (HERM/) HERMANN K.
PA (PILA/) PILARSKY C.
Query Match
Best Local Similarity 23.1%; Score 153.5; DB 8; Length 358;
RESULT 694
ID AAY41692 standard; protein; 373 AA.
DE Human PRO 363 protein sequence.
PN WO9946281-A2.
PD 16-SEP-1999.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 2; Length 373;
RESULT 695
ID AAB33430 standard; protein; 373 AA.
DE Human PRO363 protein UNQ318 SEQ ID NO:87.
PN WO200053758-A2.
PD 14-SEP-2000.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 3; Length 373;
RESULT 696
ID AAB44248 standard; protein; 373 AA.
DE Human PRO363 (UNQ318) protein sequence SEQ ID NO:59.
PN WO200053756-A2.
PD 14-SEP-2000.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 3; Length 373;
RESULT 697
ID AAU12365 standard; protein; 373 AA.
DE Human PRO363 polypeptide sequence.
PN WO200140466-A2.
PD 07-JUN-2001.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 4; Length 373;
RESULT 698
ID AAB48146 standard; protein; 373 AA.
DE Human A236 variant 2 polypeptide.
PN WO200069885-A2.
PD 23-NOV-2000.
PA (MILL-) MILLENNIUM PHARM INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 4; Length 373;
RESULT 699
ID AAB48108 standard; protein; 373 AA.
DE Human A236 polypeptide.
PN WO200069885-A2.
PD 23-NOV-2000.
PA (MILL-) MILLENNIUM PHARM INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 4; Length 373;
RESULT 700
ID AAB65293 standard; protein; 373 AA.
DE Human PRO363 protein sequence SEQ ID NO:503.
PN WO200073454-A1.
PD 07-DEC-2000.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 4; Length 373;
RESULT 701
ID AAU83656 standard; protein; 373 AA.
DE Human PRO protein, Seq ID No 130.
PN WO200208288-A2.
PD 31-JAN-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 5; Length 373;
RESULT 702
ID ABB84848 standard; protein; 373 AA.
DE Human PRO363 protein sequence SEQ ID NO:64.
PN WO200200690-A2.
PD 03-JAN-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 5; Length 373;
RESULT 703
ID AAE26448 standard; protein; 373 AA.
DE Human A236 protein.
PN US2002055139-A1.
PD 09-MAY-2002.
PA (HOLT/) HOLTZMAN D A.
PA (SHAR/) SHARP J D.
PA (LEIB/) LEIBY K R.
PA (BOSS/) BOSSONE S.
PA (PANY/) PAN Y.
PA (BARN/) BARNES T M.

PA (FRAS/) FRASER C C.
 PA (WRIG/) WRIGHTON N.
 PA (MYER/) MYERS P S.
 PA (KING/) KINGSBURY G.
 Query Match 9.1%; Score 153.5; DB 5; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 704
 ID ABB95454 standard; protein; 373 AA.
 DE Human angiogenesis related protein PRO363 SEQ ID NO: 64.
 PN W0200208284-A2.
 PD 31-JAN-2002.
 PA (GETH) GENENTECH INC.
 PA (BAKE/) BAKER K P.
 PA (FERR/) FERRARA N.
 PA (GERB/) GERBER H.
 PA (GERR/) GERRITSEN M E.
 PA (GODD/) GODDARD A.
 PA (GODO/) GODOWSKI P J.
 PA (GURN/) GURNEY A L.
 PA (HILL/) HILLAN K J.
 PA (MARS/) MARSTERS S A.
 PA (PANJ/) PAN J.
 PA (PAON/) PAONI N F.
 PA (STEP/) STEPHAN J F.
 PA (WATA/) WATANABE C K.
 PA (WILL/) WILLIAMS P M.
 PA (WOOD/) WOOD W I.
 Query Match 9.1%; Score 153.5; DB 5; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 705
 ID ABU59108 standard; protein; 373 AA.
 DE Human PRO polypeptide #140.
 PN US2003027163-A1.
 PD 06-FEB-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 6; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 706
 ID ABU59186 standard; protein; 373 AA.
 DE Novel human secreted or transmembrane protein PRO363.
 PN US2002132252-A1.
 PD 19-SEP-2002.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 6; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 707
 ID ABU82698 standard; protein; 373 AA.
 DE Human secreted/transmembrane protein PRO363.
 PN US2003032023-A1.
 PD 13-FEB-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 6; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 708
 ID ABO17809 standard; protein; 373 AA.
 DE Novel human secreted and transmembrane protein PRO363.
 PN US2003032156-A1.
 PD 13-FEB-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 6; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 709
 ID ABU60617 standard; protein; 373 AA.
 DE Human secreted/transmembrane protein, #176.
 PN US2002160384-A1.
 PD 31-OCT-2002.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 6; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 710
 ID ABU80803 standard; protein; 373 AA.
 DE Human PRO polypeptide #65.
 PN US2003036635-A1.
 PD 20-FEB-2003.
 PA (GETH) GENENTECH INC.

Query Match 9.1%; Score 153.5; DB 6; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 711
 ID ABO25194 standard; protein; 373 AA.
 DE Novel human secreted and transmembrane protein PRO363.
 PN US2003050239-A1.
 PD 13-MAR-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 6; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 712
 ID ABO33769 standard; protein; 373 AA.
 DE Novel human secreted and transmembrane protein PRO363.
 PN US2003045687-A1.
 PD 06-MAR-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 6; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 713
 ID ABU13999 standard; protein; 373 AA.
 DE Human PRO363 polypeptide.
 PN US2002103125-A1.
 PD 01-AUG-2002.
 PA (GETH) GENENTECH LTD.
 Query Match 9.1%; Score 153.5; DB 6; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 714
 ID ABU81063 standard; protein; 373 AA.
 DE Human PRO polypeptide #194.
 PN US2003004311-A1.
 PD 02-JAN-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 6; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 715
 ID ABU72200 standard; protein; 373 AA.
 DE Novel human secreted and transmembrane protein PRO363.
 PN US2002192706-A1.
 PD 19-DEC-2002.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 6; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 716
 ID ABU72584 standard; protein; 373 AA.
 DE Novel human secreted and transmembrane protein PRO363.
 PN US2003003531-A1.
 PD 02-JAN-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 6; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 717
 ID ABU66763 standard; protein; 373 AA.
 DE Human PRO polypeptide #194.
 PN US2003036180-A1.
 PD 20-FEB-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 6; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 718
 ID ABU84880 standard; protein; 373 AA.
 DE Human secreted and transmembrane polypeptide PRO363.
 PN US2002177553-A1.
 PD 28-NOV-2002.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 6; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 719
 ID ABU59844 standard; protein; 373 AA.
 DE Novel secreted and transmembrane protein PRO363.
 PN US2003017563-A1.
 PD 23-JAN-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 6; Length 373;

Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 720
ID ABU61078 standard; protein; 373 AA.
DE Human PRO363 polypeptide.
PN US2002169284-A1.
PD 14-NOV-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 721
ID ABUS9333 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, #176.
PN US2003027162-A1.
PD 06-FEB-2003.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 722
ID ABO26030 standard; protein; 373 AA.
DE Human PRO363 polypeptide.
PN US2002127576-A1.
PD 12-SEP-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 723
ID ABO25034 standard; protein; 373 AA.
DE Human secreted/transmembrane protein (PRO) #194.
PN US2003036179-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 724
ID ABUS9347 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003004102-A1.
PD 02-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 725
ID ABUS9039 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, #176.
PN US2002142961-A1.
PD 03-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 726
ID ABUS9039 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, #176.
PN US2002142961-A1.
PD 03-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 727
ID ABUS9417 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003022187-A1.
PD 30-JAN-2003.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 728
ID ABUS9482 standard; protein; 373 AA.
DE Novel human secreted or transmembrane protein PRO5723.
PN US2003027985-A1.
PD 06-FEB-2003.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 729
ID ABUS9482 standard; protein; 373 AA.
DE Human secreted/transmembrane, PRO, protein SEQ ID 388.

PN US2003032155-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 730
ID ABUS9248 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003017476-A1.
PD 23-JAN-2003.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 731
ID ABUS9248 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2002123463-A1.
PD 05-SEP-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 732
ID ABUS9248 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US200217164-A1.
PD 28-NOV-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 733
ID ABUS9248 standard; protein; 373 AA.
DE Human secreted and transmembrane polypeptide PRO363.
PN US2002197615-A1.
PD 26-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 734
ID ABUS9248 standard; protein; 373 AA.
DE Human PRO363 polypeptide.
PN US2003017981-A1.
PD 23-JAN-2003.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 735
ID ADA45907 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003022328-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 736
ID ADA45907 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003073212-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 737
ID ABUS9248 standard; protein; 373 AA.
DE Human PRO363 protein.
PN US2003050448-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Score 153.5; DB 6; Length 373;
Pred. No. 0.0002;
RESULT 738
ID ADA18988 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003054517-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.

Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 739
ID ADA61611 standard; protein; 373 AA.
DE Homo sapiens.
PN US2003049816-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 740
ID ADB19396 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003068796-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 741
ID ADB27937 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082704-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 742
ID ADA86416 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082711-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 743
ID ADB15980 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003087350-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 744
ID ADA38014 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US200308297-A1.
PD 09-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 745
ID ADA47766 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003073215-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 746
ID ADA21700 standard; protein; 373 AA.
DE Human secreted/transmembrane polypeptide PRO363.
PN US2003054404-A1.
PD 20-MAR-2003.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 747
ID ADA10487 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003059831-A1.
PD 27-MAR-2003.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 748

ID ADA67561 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003068795-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 749
ID ADB30568 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003068794-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 750
ID ADA85864 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082893-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 751
ID ADA18031 standard; protein; 373 AA.
DE Human PRO363 polypeptide.
PN US2003054987-A1.
PD 20-MAR-2003.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 752
ID ADA97076 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082705-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 753
ID ADA79380 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082763-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 754
ID ADA87519 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087345-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 755
ID ADB16721 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003087349-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 756
ID ADA28139 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003054359-A1.
PD 20-MAR-2003.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 757
ID ADA91813 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082694-A1.

PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 758
ID ADB14876 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003087451-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 759
ID ADA24598 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003050241-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 760
ID ADB18837 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003073211-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 761
ID ADA94052 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003077722-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 762
ID ADB19948 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082691-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 763
ID ADB13260 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082710-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 764
ID ABO43342 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003044945-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 765
ID ABO19649 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003050240-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 766
ID ADA1259 standard; protein; 373 AA.
DE Human secreted/transmembrane polypeptide PRO363.
PN US2003055216-A1.
PD 20-MAR-2003.

PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 767
ID ADA94719 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003059832-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 768
ID ADA74514 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003068798-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 769
ID ADB24747 standard; protein; 373 AA.
DE Human PRO polypeptide SEQ ID NO 388.
PN US2003077713-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 770
ID ADA82271 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082701-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 771
ID ADA75234 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003073216-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 772
ID ADA85312 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082695-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 773
ID ADA84760 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082708-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 774
ID ADB30016 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003073214-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 775
ID ADB80544 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082761-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 6; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 776
ID ADA1259 standard; protein; 373 AA.
DE Human secreted/transmembrane polypeptide PRO363.
PN US2003055216-A1.
PD 20-MAR-2003.


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Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 776
ID ADA75786 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082703-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 777
ID ADA38944 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003059780-A1.
PD 27-MAR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 778
ID ADA47011 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003073210-A1.
PD 17-APR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 779
ID ADB25307 standard; protein; 373 AA.
DE Human PRO polypeptide SEQ ID NO 388.
PN US2003077715-A1.
PD 24-APR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 780
ID ADA93483 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003077721-A1.
PD 24-APR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 781
ID ADB26833 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003092147-A1.
PD 15-MAY-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 782
ID ADB31120 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003096386-A1.
PD 22-MAY-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 783
ID ABJ72420 standard; protein; 373 AA.
DE Human PRO363 protein.
PN US2003027988-A1.
PD 06-FEB-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 784
ID ADA93065 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003060407-A1.
PD 27-MAR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 785
ID ADA61048 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003068797-A1.
DE Homo sapiens.
PN US2003049817-A1.
PD 13-MAR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 786
ID ADB24195 standard; protein; 373 AA.
DE Human PRO polypeptide SEQ ID NO 388.
PN US2003077714-A1.
PD 24-APR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 787
ID ADA96524 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082890-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 788
ID ADA81096 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082702-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 789
ID ADA95972 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082759-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 790
ID ADB26281 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082760-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 791
ID ADB21766 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082765-A1.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 792
ID ABO34315 standard; protein; 373 AA.
DE Human secreted/transmembrane polypeptide PRO 363.
PN US2003044934-A1.
PD 06-MAR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 793
ID ABO19540 standard; protein; 373 AA.
DE Novel human secreted and transmembrane polypeptide #8.
PN US2003049633-A1.
PD 13-MAR-2003.
PA (GETH ) GENENTECH INC. 9.1%; Score 153.5; DB 6; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 794
ID ADA77545 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003068797-A1.
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PD 10-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 795
ID ADB18285 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003077710-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 796
ID ADA86968 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082709-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 797
ID ADA88071 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082700-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 798
ID ADA46459 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003054516-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 799
ID ADB28489 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082699-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 800
ID ADB29041 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082706-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 801
ID ABO53245 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003044806-A1.
PD 06-MAR-2003.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 802
ID ADA76993 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003059909-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 803
ID ADA22626 standard; protein; 373 AA.
DE Human secreted/transmembrane polypeptide PRO363.
PN US2003040473-A1.
PD 27-FEB-2003.
Query Match 9.1%; Score 153.5; DB 7; Length 373;

Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 804
ID ADA8623 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003073213-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 805
ID ADA97628 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082686-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 806
ID ADB27385 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003022239-A1.
PD 30-JAN-2003.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 807
ID ADB22318 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087344-A1.
PD 08-MAY-2003.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 808
ID ABO22615 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003017982-A1.
PD 23-JAN-2003.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 809
ID ADA06792 standard; protein; 373 AA.
DE Human secreted/transmembrane PRO polypeptide #140.
PN US2003049638-A1.
PD 13-MAR-2003.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 810
ID ABJ72122 standard; protein; 373 AA.
DE Human membrane bound receptor/protein PRO363 amino acid sequence.
PN US2003065147-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 811
ID ADA39485 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003059782-A1.
PD 27-MAR-2003.
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 812
ID ADA67009 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003068793-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 813
ID ADB22870 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003077711-A1.
PD 24-APR-2003.

PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 814
ID ADB23643 standard; protein; 373 AA.
DE Human PRO polypeptide SEQ ID NO 388.
PN US2003077712-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 815
ID ADA92365 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082712-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 816
ID ADB15428 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003087352-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 817
ID ADB83620 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003073814-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 818
ID ADB80726 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003088068-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 819
ID ADB73267 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003096968-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 820
ID ADB38680 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082766-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 821
ID ADB96511 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003054403-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 822
ID ADB78349 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092889-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 823
ID ADB38128 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087347-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 824
ID ADB66600 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082689-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 825
ID ADB84997 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003073817-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 826
ID ADB89680 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082698-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 827
ID ADB90412 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003082762-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 828
ID ADB39513 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082764-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 829
ID ADB78103 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092886-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 830
ID ADB73565 standard; protein; 373 AA.
DE Human PRO polypeptide #8.
PN US2003045462-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 831
ID ADB87169 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003088067-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;

ID ADB33953 standard; protein; 373 AA.
DE Human PRO polypeptide SEQ ID NO 388.
FN US2003077716-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Similarity 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 842
ID ADB33057 standard; protein; 373 AA.
DE Human PRO polypeptide SEQ ID NO 388.
FN US2003077718-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Similarity 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 843
ID ADB36161 standard; protein; 373 AA.
DE Human PRO polypeptide SEQ ID NO 388.
FN US2003077720-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Similarity 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 844
ID ADB46556 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
FN US2003082692-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Similarity 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 845
ID ADC43707 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
FN US2003054986-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Similarity 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 846
ID ADC57983 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
FN US2003027754-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Similarity 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 847
ID ADC55347 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
FN US2003045463-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Similarity 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 848
ID ADC12214 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
FN US2003049681-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Similarity 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 849
ID ADC61467 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
FN US2003049684-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Similarity 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 850
ID ADC63431 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
FN US2003054405-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Similarity 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;

PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 851
ID ADC66531 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003060406-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 852
ID ADC56636 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003064375-A1.
PD 03-APR-2003.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 853
ID ADC68655 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003064407-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 854
ID ADC62715 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003068648-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 855
ID ADC67780 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003069178-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 856
ID ADC11681 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003069403-A1.
PD 10-APR-2003.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 857
ID ADC41100 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003072745-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 858
ID ADC67155 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003073131-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 859
ID ADC62091 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003073624-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 860
ID ADC36859 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US200308065-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 861
ID ADC41724 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003104998-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 862
ID ADC21849 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003096969-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 863
ID ADC50429 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092106-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 864
ID ADC71976 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092107-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 865
ID ADC59955 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092105-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 866
ID ADC49880 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003088064-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 867
ID ADC49079 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003088070-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 868
ID ADC49596 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003088071-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 869
ID ADC49596 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003088071-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 27.0%; Pred. No. 0.0002;

ID ADC47457 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003088072-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 870
ID ADC52962 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein Seq ID388.
PN US2003087365-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 871
ID ADC57316 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein Seq ID388.
PN US2003087366-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 872
ID ADC60507 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087367-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 873
ID ADC50982 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087361-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 874
ID ADC65509 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003087362-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 875
ID ADC54607 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein Seq ID388.
PN US2003087363-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 876
ID ADC53568 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein Seq ID388.
PN US2003087364-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 877
ID ADC59091 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein Seq ID388.
PN US2003087359-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 878
ID ADC55969 standard; protein; 373 AA.

DE Novel human secreted and transmembrane protein Seq ID388.
PN US2003087360-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 879
ID ADC58539 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein Seq ID388.
PN US2003087346-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 880
ID ADC14803 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003082546-A1.
PD 01-MAY-2003.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 881
ID ADC47202 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003105288-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 882
ID ADD08335 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003068623-A1.
PD 10-APR-2003.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 883
ID ADD03213 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092104-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 884
ID ADC90205 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087348-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 885
ID ADC82160 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003083461-A1.
PD 01-MAY-2003.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 886
ID ADC69624 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194770-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 887
ID ADC48513 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194773-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.

Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 888
ID ADC47961 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194771-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 889
ID ADD07802 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2002193299-A1.
PD 19-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 890
ID ADC78077 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003096972-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 891
ID ADD04617 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087354-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 892
ID ADC82693 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003059833-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 893
ID ADD06312 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003073816-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 894
ID ADC80573 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092103-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 895
ID ADD11080 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194774-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 896
ID ADD10353 standard; protein; 373 AA.
DE Human secreted/transmembrane PRO polypeptide #32.
PN US2003105011-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;

RESULT 897
ID ADC47961 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194771-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 898
ID ADD08873 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003073090-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 899
ID ADC77831 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003088066-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 900
ID ADC80021 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087358-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 901
ID ADD07122 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2002193300-A1.
PD 19-DEC-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 902
ID ADD11313 standard; protein; 373 AA.
DE Human secreted/transmembrane PRO polypeptide #32.
PN US2003105013-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 903
ID ADD09490 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194775-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 904
ID ADC83369 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003059783-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 905
ID ADD50794 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003105291-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 906
ID ADD41203 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.

PN US2003203438-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 907
ID ADD52342 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194769-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 908
ID ADD51040 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003105290-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 909
ID ADD53082 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194732-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 910
ID ADD53634 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003203437-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 911
ID ADD5476 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003077593-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 912
ID ADD37106 standard; protein; 373 AA.
DE Human secreted/transmembrane PRO polypeptide #32.
PN US2003105012-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 913
ID ADD56434 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003077594-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 914
ID ADD51790 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194779-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 915
ID ADD02589 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003203431-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.

Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 916
ID ADD50521 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003096971-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 917
ID ADD02023 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003203430-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 918
ID ADD54205 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003203432-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 919
ID ADD54872 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2002132253-A1.
PD 19-SEP-2002.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 920
ID ADD50275 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003096970-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 921
ID ADD51286 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003105289-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 922
ID ADD49093 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003096744-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 923
ID ADD92522 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199030-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 924
ID ADD91418 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199055-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;


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Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 925
ID ADE04032 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199057-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 926
ID ADE31891 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US200308647-A1.
PD 10-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 927
ID ADE27026 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087304-A1.
PD 08-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 928
ID ADE32329 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003194765-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 929
ID ADE22261 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199056-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 930
ID ADD79485 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003203428-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 931
ID ADE35147 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003203434-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 932
ID ADE16261 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003203435-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 933
ID ADD72876 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003203436-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 934
ID ADE42021 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194772-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 935
ID ADE17838 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199023-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 936
ID ADD91970 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199053-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 937
ID ADE33433 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003194767-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 938
ID ADE33985 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003194791-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 939
ID ADD80037 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207417-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 940
ID ADD93074 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194768-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 941
ID ADD72234 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003194781-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 942
ID ADE19494 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199025-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 943
ID ADE18942 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
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PN US2003199026-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 944
ID ADE43138 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199033-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 945
ID ADD95927 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199059-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 946
ID ADE22813 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199064-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 947
ID ADD78931 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003203429-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 948
ID ADE26493 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087305-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 949
ID ADE32881 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003194766-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 950
ID ADE42573 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199032-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 951
ID ADE16885 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003203433-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 952
ID ADD80589 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207418-A1.
PD 06-NOV-2003.

PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 953
ID AD89617 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199028-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 954
ID ADE40901 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199031-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 955
ID ADE04700 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199034-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 956
ID ADE92829 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194777-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 957
ID ADF46899 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003195333-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 958
ID ADF67430 standard; protein; 373 AA.
DE Human PRO363 amino acid sequence SEQ ID NO:503.
PN US2002198148-A1.
PD 26-DEC-2002.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 959
ID ADG21538 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207355-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 960
ID ADG23179 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207384-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match 27.0%; Pred. No. 0.0002;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 961
ID ADF97514 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207370-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.

Query Match 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 962
ID ADG80578 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207373-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 963
ID ADG52656 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003216561-A1.
PD 20-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 964
ID ADG59976 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003206915-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 965
ID ADG80026 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207372-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 966
ID ADH55318 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207381-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 967
ID ADH55870 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207379-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 968
ID ADI35684 standard; protein; 373 AA.
DE Human PRO polypeptide #140.
PN US2003050457-A1.
PD 13-MAR-2003.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 969
ID ADI60736 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003077000-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 970
ID ADI64089 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207385-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;

RESULT 971
ID ADI65038 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207386-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 972
ID ADI63537 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207387-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 973
ID ADH81951 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207388-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 974
ID ADI00177 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003049682-A1.
PD 13-MAR-2003.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 975
ID ADH81399 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207377-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 976
ID ADM82568 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087355-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 977
ID ADN15967 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087353-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 978
ID ADN16596 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087385-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 979
ID ADN15415 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003087356-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 7; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 980
ID ADN14863 standard; protein; 373 AA.

DE Novel human secreted and transmembrane protein PRO363.
PN US2003087357-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 7; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 981
ID ADC48833 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092888-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 982
ID ADC81125 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003092115-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 983
ID ADE21004 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100735-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 984
ID ADE05848 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100728-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 985
ID ADD76573 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003100087-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 986
ID ADD75077 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100712-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 987
ID ADD75823 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100717-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 988
ID ADD85055 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100722-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 989
ID ADD86881 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.

PN US2003100738-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 990
ID ADE20758 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100734-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 991
ID ADE39055 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003096362-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 992
ID ADD87937 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003092113-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 993
ID ADD86341 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003203440-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 994
ID ADE05602 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100727-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 995
ID ADD73587 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100711-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 996
ID ADE75789 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003211571-A1.
PD 13-NOV-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 997
ID ADE48393 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003104536-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC. 9.1%; Score 153.5; DB 8; Length 373;
Query Match Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 998
ID ADD78427 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100737-A1.

PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 999
ID ADE41314 standard; protein; 373 AA.
DE Human secreted/transmembrane PRO polypeptide #32.
PN US2003100497-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1000
ID ADE23365 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003092108-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1001
ID ADE21250 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100736-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1002
ID ADD77365 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100732-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1003
ID ADE20512 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100733-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1004
ID ADD75577 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100084-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1005
ID ADD74093 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100708-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1006
ID ADD74339 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100709-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1007
ID ADD76069 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100718-A1.
PD 29-MAY-2003.

PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1008
ID ADD85561 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100721-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1009
ID ADE23917 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003092110-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1010
ID ADE24560 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003092111-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1011
ID ADD87385 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003203439-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1012
ID ADE05110 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100726-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1013
ID ADD75323 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003100714-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1014
ID ADD76867 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100715-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1015
ID ADD86635 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003100719-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1016
ID ADE89251 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199062-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.

Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1017
 ID ADD78103 standard; protein; 373 AA.
 DE Novel human secreted and transmembrane protein PRO363.
 PN US2003100731-A1.
 PD 29-MAY-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1018
 ID ADE18390 standard; protein; 373 AA.
 DE Human PRO polypeptide #194.
 PN US2003194794-A1.
 PD 16-OCT-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1019
 ID ADE86699 standard; protein; 373 AA.
 DE Human PRO polypeptide #194.
 PN US2003199054-A1.
 PD 23-OCT-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1020
 ID ADE89494 standard; protein; 373 AA.
 DE Human secreted/transmembrane protein, PRO363.
 PN US2003130181-A1.
 PD 10-JUL-2003.
 PA (ASHK/) ASHKENAZI A J.
 PA (BAKE/) BAKER K P.
 PA (BOTS/) BOTSTEIN D.
 PA (DESN/) DESNOYERS L.
 PA (EATO/) EATON D L.
 PA (FERR/) FERRARA N.
 PA (FILV/) FILVAROFF E.
 PA (FONG/) FONG S.
 PA (GAOW/) GAO W.
 PA (GERB/) GERBER H.
 PA (GERR/) GERRITSEN M E.
 PA (GODD/) GODDARD A.
 PA (GODO/) GODOWSKI P J.
 PA (GIRM/) GIRMALDI J C.
 PA (GURN/) GURNEY A L.
 PA (HILL/) HILLAN K J.
 PA (KLJA/) KLJAVIN I J.
 PA (KUOS/) KUO S S.
 PA (NAPI/) NAPIER M A.
 PA (PANJ/) PAN J.
 PA (PAON/) PAONI N F.
 PA (ROYM/) ROY M A.
 PA (SHEL/) SHELTON D L.
 PA (STEW/) STEWART T A.
 PA (TUMA/) TUMAS D.
 PA (WILL/) WILLIAMS P M.
 PA (WOOD/) WOOD W I.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1021
 ID ADD77611 standard; protein; 373 AA.
 DE Novel human secreted and transmembrane protein PRO363.
 PN US2003100729-A1.
 PD 29-MAY-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1022
 ID ADD77857 standard; protein; 373 AA.
 DE Novel human secreted and transmembrane protein PRO363.
 PN US2003100730-A1.
 PD 29-MAY-2003.
 PA (GETH) GENENTECH INC.

PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1023
 ID ADD85315 standard; protein; 373 AA.
 DE Novel human secreted and transmembrane protein PRO363.
 PN US2003100725-A1.
 PD 29-MAY-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1024
 ID ADD73847 standard; protein; 373 AA.
 DE Human PRO polypeptide #65.
 PN US2003100710-A1.
 PD 29-MAY-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1025
 ID ADD74585 standard; protein; 373 AA.
 DE Human PRO polypeptide #65.
 PN US2003100713-A1.
 PD 29-MAY-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1026
 ID ADD77113 standard; protein; 373 AA.
 DE Novel human secreted and transmembrane protein PRO363.
 PN US2003100716-A1.
 PD 29-MAY-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1027
 ID ADD85807 standard; protein; 373 AA.
 DE Novel human secreted and transmembrane protein PRO363.
 PN US2003100720-A1.
 PD 29-MAY-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1028
 ID ADE05356 standard; protein; 373 AA.
 DE Human PRO polypeptide #65.
 PN US2003100723-A1.
 PD 29-MAY-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1029
 ID ADD74831 standard; protein; 373 AA.
 DE Human PRO polypeptide #65.
 PN US2003100724-A1.
 PD 29-MAY-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1030
 ID ADF61134 standard; protein; 373 AA.
 DE Human secreted/transmembrane protein, PRO363.
 PN US2003195345-A1.
 PD 16-OCT-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1031
 ID ADF39826 standard; protein; 373 AA.
 DE Human secreted/transmembrane protein, PRO363.
 PN US2003198994-A1.
 PD 23-OCT-2003.
 PA (GETH) GENENTECH INC.

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Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1032
ID ADF45622 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003195148-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1033
ID ADE94719 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199027-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1034
ID ADE91130 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199061-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1035
ID ADF35629 standard; protein; 373 AA.
DE Human PRO363 polypeptide.
PN US2003194760-A1.
PD 16-OCT-2003.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1036
ID ADE95271 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199052-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1037
ID ADE93381 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199060-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1038
ID ADF24018 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003204055-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1039
ID ADF40450 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003199021-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1040
ID ADF23394 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003203402-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1041
ID ADF33377 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003194780-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1042
ID ADF34962 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199029-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1043
ID ADF26844 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003199436-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1044
ID ADF27480 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003199437-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1045
ID ADE92277 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003199051-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1046
ID ADE90578 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003199063-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1047
ID ADF41074 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003199435-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1048
ID ADF32753 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003211091-A1.
PD 13-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1049
ID ADF25119 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003211092-A1.
PD 13-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match          9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1050
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ID ADF26220 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003199674-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1051 27.0%; Pred. No. 0.0002;
ID ADF34009 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003194410-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1052 27.0%; Pred. No. 0.0002;
ID ADF46246 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003195344-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1053 27.0%; Pred. No. 0.0002;
ID ADE91725 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003199058-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1054 27.0%; Pred. No. 0.0002;
ID ADG11879 standard; protein; 373 AA.
DE Human PRO363 polypeptide.
PN US2003228655-A1.
PD 11-DEC-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1055 27.0%; Pred. No. 0.0002;
ID ADG05643 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003096959-A1.
PD 22-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1056 27.0%; Pred. No. 0.0002;
ID ADG27197 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003096962-A1.
PD 22-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1057 27.0%; Pred. No. 0.0002;
ID ADG02304 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207352-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1058 27.0%; Pred. No. 0.0002;
ID ADG22090 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207360-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1059 27.0%; Pred. No. 0.0002;
ID ADG20160 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207376-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1060 27.0%; Pred. No. 0.0002;
ID ADF98066 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207422-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1061 27.0%; Pred. No. 0.0002;
ID ADG24283 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207426-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1062 27.0%; Pred. No. 0.0002;
ID ADF98637 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003208055-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1063 27.0%; Pred. No. 0.0002;
ID ADG03468 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207351-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1064 27.0%; Pred. No. 0.0002;
ID ADF99189 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207353-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1065 27.0%; Pred. No. 0.0002;
ID ADG16774 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207359-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1066 27.0%; Pred. No. 0.0002;
ID ADG05233 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207375-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1067 27.0%; Pred. No. 0.0002;
ID ADG19500 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207425-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9.1%; Score 153.5; DB 8; Length 373;
RESULT 1068 27.0%; Pred. No. 0.0002;
ID ADG11260 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
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PN US2003096967-A1.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1069
ID ADG13337 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207357-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1070
ID ADG08394 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207424-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1071
ID ADG15564 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003219885-A1.
PD 27-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1072
ID ADG12039 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003096963-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1073
ID ADF96962 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207371-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1074
ID ADG06147 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207374-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1075
ID ADG23731 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207389-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1076
ID ADG04020 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207423-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1077
ID ADG24921 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207427-A1.

PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1078
ID ADF94596 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003096964-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1079
ID ADG07218 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207350-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1080
ID ADG07770 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207356-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1081
ID ADG06932 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003096966-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1082
ID ADG55265 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003194778-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1083
ID ADG60929 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207390-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1084
ID ADG62033 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207428-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1085
ID ADG82234 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207358-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1086
ID ADG57473 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207362-A1.
PD 06-NOV-2003.

Query Match	9.1%;	Score 153.5;	DB 8;	Length 373;
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1096				
ID ADG81682 standard; protein;	373 AA.			
DE Human PRO polypeptide #194.				
FN US2003207805-A1.				
PD 06-NOV-2003.				
Query Match	9.1%;	Score 153.5;	DB 8;	Length 373;
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1097				
ID ADH19749 standard; protein;	373 AA.			
DE Human secreted/transmembrane protein PRO363.				
FN US2003228656-A1.				
PD 11-DEC-2003.				
PA (GETH) GENENTECH INC.				
Query Match	9.1%;	Score 153.5;	DB 8;	Length 373;
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1098				
ID ADH30644 standard; protein;	373 AA.			
DE Human PRO polypeptide #194.				
FN US2003077723-A1.				
PD 24-APR-2003.				
PA (GETH) GENENTECH INC.				
Query Match	9.1%;	Score 153.5;	DB 8;	Length 373;
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1099				
ID ADH12011 standard; protein;	373 AA.			
DE Novel human secreted and transmembrane protein PRO363.				
FN US2003207419-A1.				
PD 06-NOV-2003.				
PA (GETH) GENENTECH INC.				
Query Match	9.1%;	Score 153.5;	DB 8;	Length 373;
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1100				
ID ADG49608 standard; protein;	373 AA.			
DE Human secreted/transmembrane protein, PRO363.				
FN US2003215905-A1.				
PD 20-NOV-2003.				
PA (GETH) GENENTECH INC.				
Query Match	9.1%;	Score 153.5;	DB 8;	Length 373;
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1101				
ID ADG51480 standard; protein;	373 AA.			
DE Human secreted/transmembrane protein, PRO363.				
FN US2003215908-A1.				
PD 20-NOV-2003.				
PA (GETH) GENENTECH INC.				
Query Match	9.1%;	Score 153.5;	DB 8;	Length 373;
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1102				
ID ADG52433 standard; protein;	373 AA.			
DE Novel human secreted and transmembrane protein PRO363.				
FN US2003207414-A1.				
PD 06-NOV-2003.				
PA (GETH) GENENTECH INC.				
Query Match	9.1%;	Score 153.5;	DB 8;	Length 373;
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1103				
ID ADG54161 standard; protein;	373 AA.			
DE Novel human secreted and transmembrane protein PRO363.				
FN US2003207416-A1.				
PD 06-NOV-2003.				
PA (GETH) GENENTECH INC.				
Query Match	9.1%;	Score 153.5;	DB 8;	Length 373;
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1104				
ID ADG48984 standard; protein;	373 AA.			
DE Human secreted/transmembrane protein, PRO363.				
FN US2003216305-A1.				
PD 20-NOV-2003.				
PA (GETH) GENENTECH INC.				
Query Match	9.1%;	Score 153.5;	DB 8;	Length 373;
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1105				
ID ADG54161 standard; protein;	373 AA.			
DE Novel human secreted and transmembrane protein PRO363.				
FN US2003207416-A1.				
PD 06-NOV-2003.				
PA (GETH) GENENTECH INC.				
Query Match	9.1%;	Score 153.5;	DB 8;	Length 373;
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1106				
ID ADG48984 standard; protein;	373 AA.			
DE Human secreted/transmembrane protein, PRO363.				
FN US2003216305-A1.				
PD 20-NOV-2003.				
PA (GETH) GENENTECH INC.				
Query Match	9.1%;	Score 153.5;	DB 8;	Length 373;
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1107				
ID ADG48984 standard; protein;	373 AA.			
DE Human secreted/transmembrane protein, PRO363.				
FN US2003216305-A1.				
PD 20-NOV-2003.				
PA (GETH) GENENTECH INC.				
Query Match	9.1%;	Score 153.5;	DB 8;	Length 373;
Best Local Similarity	27.0%;	Pred. No. 0.0002;		
RESULT 1108				
ID ADG48984 standard; protein;	373 AA.			
DE Human secreted/transmembrane protein, PRO363.				
FN US2003216305-A1.				
PD 20-NOV-2003.				
PA (GETH) GENENTECH INC.				
Query Match	9.1%;	Score 15		

RESULT 1105
ID ADG81130 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003194793-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1106
ID ADG56369 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207366-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1107
ID ADH12635 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207378-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1108
ID ADG48360 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2003216560-A1.
PD 20-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1109
ID ADH21242 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003224358-A1.
PD 04-DEC-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1110
ID ADG61481 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207429-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1111
ID ADH20282 standard; protein; 373 AA.
DE Human secreted/transmembrane protein PRO363.
PN US2003219856-A1.
PD 27-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1112
ID ADH28568 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003022331-A1.
PD 30-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1113
ID ADG54713 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207367-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1114
ID ADG59753 standard; protein; 373 AA.

DE Novel human secreted and transmembrane protein PRO363.
PN US2003207369-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1115
ID ADG50856 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2004005312-A1.
PD 08-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1116
ID ADH43497 standard; protein; 373 AA.
DE Human PRO polypeptide #32.
PN US2003224984-A1.
PD 04-DEC-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1117
ID ADG58800 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2004005657-A1.
PD 08-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1118
ID ADG34126 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2004006206-A1.
PD 08-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1119
ID ADG62256 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2004006219-A1.
PD 08-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1120
ID ADI81177 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2003207361-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1121
ID ADI33596 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2003096960-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1122
ID ADH69690 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2004019183-A1.
PD 29-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1123
ID ADH25281 standard; protein; 373 AA.
DE Human neurotrophin homologue related protein sequence SEQ ID NO:59.

PN EPI386931-A1.
PD 04-FEB-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1124
ID ADG09920 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2004009548-A1.
PD 15-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1125
ID ADI15391 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207382-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1126
ID ADG09268 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2004009547-A1.
PD 15-JAN-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1127
ID ADI14723 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207383-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1128
ID ADI29851 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003096961-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1129
ID ADI18318 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2003207349-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1130
ID ADM27248 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2004044179-A1.
PD 04-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1131
ID ADJ63599 standard; protein; 373 AA.
DE Novel human secreted and transmembrane protein PRO363.
PN US2004039164-A1.
PD 26-FEB-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1132
ID ADJ77494 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2004038336-A1.

PD 26-FEB-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1133
ID ADK82842 standard; protein; 373 AA.
DE Human PRO polypeptide #32.
PN US2004043927-A1.
PD 04-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1134
ID ADK66606 standard; protein; 373 AA.
DE Human PRO polypeptide #65.
PN US2004044180-A1.
PD 04-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1135
ID ADJ65616 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2004038335-A1.
PD 26-FEB-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1136
ID ADM27752 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2004048333-A1.
PD 11-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1137
ID ADM17058 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2004048332-A1.
PD 11-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1138
ID ADJ06892 standard; protein; 373 AA.
DE Human secreted/transmembrane protein, PRO363.
PN US2004063921-A1.
PD 01-APR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1139
ID ADM42476 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2004058424-A1.
PD 25-MAR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1140
ID ADN05373 standard; protein; 373 AA.
DE Antipsoriatic protein sequence #858.
PN WO2004028479-A2.
PD 08-APR-2004.
PA (GETH) GENENTECH INC.
Query Match 9.1%; Score 153.5; DB 8; Length 373;
Best Local Similarity 27.0%; Pred. No. 0.0002;
RESULT 1141
ID ADM28338 standard; protein; 373 AA.
DE Human PRO polypeptide #194.
PN US2004077064-A1.
PD 22-APR-2004.

PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1142
 ID ADO36720 standard; protein; 373 AA.
 DE Human UKW polypeptide, SEQ ID NO:2.
 PN EP1416279-A1.
 PD 06-MAY-2004.
 PA (HOFF) HOFFMANN LA ROCHE & CO AG F.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1143
 ID ADI95820 standard; protein; 373 AA.
 DE Human PRO polypeptide #194.
 PN US2003077659-A1.
 PD 24-APR-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1144
 ID ADI96372 standard; protein; 373 AA.
 DE Novel human secreted and transmembrane protein PRO363.
 PN US2003207354-A1.
 PD 06-NOV-2003.
 PA (GETH) GENENTECH INC.
 Query Match 9.1%; Score 153.5; DB 8; Length 373;
 Best Local Similarity 27.0%; Pred. No. 0.0002;
 RESULT 1145
 ID ADA54925 standard; protein; 512 AA.
 DE Human protein, SEQ ID 2493.
 PN EP1293569-A2.
 PD 19-MAR-2003.
 PA (HELI-) HELIX RES INST.
 Query Match 9.1%; Score 153.5; DB 6; Length 512;
 Best Local Similarity 25.2%; Pred. No. 0.00031;
 RESULT 1146
 ID AAY53666 standard; protein; 4412 AA.
 DE Sequence gi/1017427/emb/CAAG2189 from an alignment with protein 608.
 PN WO9960164-A1.
 PD 25-NOV-1999.
 PA (QUAR-) QUARK BIOTECH INC.
 Query Match 9.1%; Score 153.5; DB 3; Length 4412;
 Best Local Similarity 22.0%; Pred. No. 0.0058;
 RESULT 1147
 ID ABP60991 standard; protein; 5635 AA.
 DE Novel human protein. SEQ ID 78.
 PN WO200250105-A1.
 PD 27-JUN-2002.
 PA (SMIK) SMITHKLINE BEECHAM CORP.
 PA (SMIK) SMITHKLINE BEECHAM PLC.
 PA (GLAX) GLAXO GROUP LTD.
 Query Match 9.1%; Score 153.5; DB 5; Length 5635;
 Best Local Similarity 25.2%; Pred. No. 0.0081;
 RESULT 1148
 ID ADJ70089 standard; protein; 5636 AA.
 DE Human heat mitochondrial protein as a therapeutic target SeqID1895.
 PN WO2003087768-A2.
 PD 23-OCT-2003.
 PA (MITO-) MITOKOR.
 PA (BUCK-) BUCK INST AGE RES.
 Query Match 9.1%; Score 153.5; DB 7; Length 5636;
 Best Local Similarity 25.2%; Pred. No. 0.0081;
 RESULT 1149
 ID ADJ83137 standard; protein; 5636 AA.
 DE Human hemectin protein - SEQ ID 128.
 PN US2003170630-A1.
 PD 11-SEP-2003.
 PA (ALSO/) ALSOBROOK J P.
 PA (TCHE/) TCHERNEV V T.
 PA (LIUX/) LIU X.
 PA (SPYT/) SPYTEK K A.
 PA (ZERR/) ZERRHUSEN B D.

PA (PATT/) PATTURAJAN M.
 PA (LEPL/) LEPLEY D M.
 PA (BURG/) BURGESS C E.
 PA (SHIM/) SHIMKETS R A.
 PA (GROS/) GROSSE W M.
 PA (SZEK/) SZEKERES E S.
 PA (VERN/) VERNET C A M.
 PA (LILL/) LI L.
 PA (CASM/) CASMAN S J.
 PA (BOLD/) BOLDOG F L.
 PA (GORM/) GORMAN L.
 PA (GANG/) GANGOLLI E A.
 PA (FERN/) FERNANDES E R.
 PA (RIEG/) RIEGER D K.
 PA (EDIN/) EDINGER S R.
 PA (GUNT/) GUNTHER E.
 PA (MILL/) MILLET I.
 PA (SCIO/) SCIORE P.
 PA (ELLE/) ELLERMAN K.
 PA (MACD/) MACDOUGALL J R.
 PA (SMIT/) SMITHSON G.
 Query Match 9.1%; Score 153.5; DB 7; Length 5636;
 Best Local Similarity 25.2%; Pred. No. 0.0081;
 RESULT 1150
 ID ADK60205 standard; protein; 5636 AA.
 DE Angiogenesis differentially expressed protein GS-P29.
 PN FR2836687-A1.
 PD 05-SEP-2003.
 PA (GENE-) GENE SIGNAL.
 PA (ALMA/) AL MAHMOOD S.
 Query Match 9.1%; Score 153.5; DB 8; Length 5636;
 Best Local Similarity 25.2%; Pred. No. 0.0081;
 RESULT 1151
 ID ADK60506 standard; protein; 5636 AA.
 DE Angiogenesis differentially expressed protein GS-P29.
 PN FR2836686-A1.
 PD 05-SEP-2003.
 PA (GENE-) GENE SIGNAL.
 PA (ALMA/) AL MAHMOOD S.
 Query Match 9.1%; Score 153.5; DB 8; Length 5636;
 Best Local Similarity 25.2%; Pred. No. 0.0081;
 RESULT 1152
 ID ADP73129 standard; protein; 5636 AA.
 DE Angiogenesis inhibitor human protein sequence, GS-P29.
 PN FR2843753-A1.
 PD 27-FEB-2004.
 PA (GENE/) GENE S.
 PA (ALMS/) AL M S.
 Query Match 9.1%; Score 153.5; DB 8; Length 5636;
 Best Local Similarity 25.2%; Pred. No. 0.0081;
 RESULT 1153
 ID ADA50158 standard; protein; 393 AA.
 DE Human CAR/mouse SCF mature fusion protein.
 PN US2003092068-A1.
 PD 15-MAY-2003.
 PA (ITOH/) ITOH A.
 PA (HANA/) HANAZONO Y.
 PA (OKAD/) OKADA T.
 PA (OZAW/) OZAWA K.
 Query Match 9.1%; Score 153; DB 6; Length 393;
 Best Local Similarity 24.3%; Pred. No. 0.00024;
 RESULT 1154
 ID ABG74786 standard; protein; 31267 AA.
 DE Human RGS11 protein.
 PN WO2002103355-A1.
 PD 27-DEC-2002.
 PA (TAKE) TAKEDA CHEM IND LTD.
 Query Match 9.1%; Score 153; DB 6; Length 31267;
 Best Local Similarity 28.2%; Pred. No. 0.091;
 RESULT 1155
 ID ADA50159 standard; protein; 474 AA.
 DE Human CAR/mouse anti-CD34 antibody mature fusion protein.
 PN US2003092068-A1.

PD 15-MAY-2003.
PA (ITOH/) ITOH A.
PA (HANA/) HANAZONO Y.
PA (OKAD/) OKADA T.
PA (OZAW/) OZAWA K.
Query Match
Best Local Similarity 9.0%; Score 152; DB 6; Length 474;
23.2%; Pred. No. 0.00038;
RESULT 1156
ID AAB48145 standard; protein; 373 AA.
DE Human A236 variant 1 polypeptide.
PN WO200069885-A2.
PD 23-NOV-2000.
PA (MILL-) MILLENNIUM PHARM INC.
Query Match
Best Local Similarity 9.0%; Score 151.5; DB 4; Length 373;
27.0%; Pred. No. 0.0003;
RESULT 1157
ID AAU17996 standard; protein; 301 AA.
DE Human immunoglobulin polypeptide SEQ ID NO 141.
PN WO20015315-A2.
PD 02-AUG-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match
Best Local Similarity 8.9%; Score 151; DB 4; Length 301;
25.8%; Pred. No. 0.00025;
RESULT 1158
ID ABB10232 standard; protein; 301 AA.
DE Human cDNA SEQ ID NO: 540.
PN WO200154474-A2.
PD 02-AUG-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match
Best Local Similarity 8.9%; Score 151; DB 4; Length 301;
25.8%; Pred. No. 0.00025;
RESULT 1159
ID ABP66819 standard; protein; 301 AA.
DE Human polypeptide SEQ ID NO 540.
PN US2002090672-A1.
PD 11-JUL-2002.
PA (ROSE/) ROSEN C A.
PA (RUBE/) RUBEN S M.
PA (BARA/) BARASH S C.
Query Match
Best Local Similarity 8.9%; Score 151; DB 5; Length 301;
25.8%; Pred. No. 0.00025;
RESULT 1160
ID ADB31620 standard; protein; 301 AA.
DE Human novel protein SEQ ID NO 141.
PN US2003077606-A1.
PD 24-APR-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match
Best Local Similarity 8.9%; Score 151; DB 7; Length 301;
25.8%; Pred. No. 0.00025;
RESULT 1161
ID ADR41522 standard; protein; 318 AA.
DE Human CD-like molecule HKAC103, SEQ ID NO:321.
PN WO200226930-A2.
PD 04-APR-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match
Best Local Similarity 8.9%; Score 151; DB 5; Length 318;
25.8%; Pred. No. 0.00027;
RESULT 1162
ID ADA50157 standard; protein; 393 AA.
DE Human CAR/SCF mature fusion protein.
PN US2003092068-A1.
PD 15-MAY-2003.
PA (ITOH/) ITOH A.
PA (HANA/) HANAZONO Y.
PA (OKAD/) OKADA T.
PA (OZAW/) OZAWA K.
Query Match
Best Local Similarity 8.9%; Score 151; DB 6; Length 393;
25.4%; Pred. No. 0.00036;
RESULT 1163
ID ABG02117 standard; protein; 434 AA.
DE Novel human diagnostic protein #2108.
PN WO200175067-A2.
PD 11-OCT-2001.

PA (HYSE-) HYSEQ INC.
Query Match
Best Local Similarity 8.9%; Score 151; DB 4; Length 434;
24.3%; Pred. No. 0.00041;
RESULT 1164
ID ABU62399 standard; protein; 466 AA.
DE Chimeric CAR/Hg/Pro-A gene product.
PN US6524572-B1.
PD 25-FEB-2003.
PA (RAIN-) RAINBOW THERAPEUTIC CO.
Query Match
Best Local Similarity 8.9%; Score 151; DB 6; Length 466;
25.4%; Pred. No. 0.00045;
RESULT 1165
ID AAB48147 standard; protein; 373 AA.
DE Human A236 variant 3 polypeptide.
PN WO200069885-A2.
PD 23-NOV-2000.
PA (MILL-) MILLENNIUM PHARM INC.
Query Match
Best Local Similarity 8.9%; Score 150.5; DB 4; Length 373;
26.6%; Pred. No. 0.00037;
RESULT 1166
ID ABG31321 standard; protein; 2652 AA.
DE Predicted human adican-2 protein.
PN WO200246364-A2.
PD 13-JUN-2002.
PA (QUAR-) QUARK BIOTECH INC.
Query Match
Best Local Similarity 8.7%; Score 147.5; DB 5; Length 2652;
21.8%; Pred. No. 0.0095;
RESULT 1167
ID ADU02250 standard; protein; 2652 AA.
DE Human OCP protein #6.
PN US2004053301-A1.
PD 18-MAR-2004.
PA (QUAR-) QUARK BIOTECH INC.
Query Match
Best Local Similarity 8.7%; Score 147.5; DB 8; Length 2652;
21.8%; Pred. No. 0.0095;
RESULT 1168
ID ABP70049 standard; protein; 2845 AA.
DE Human NOVIA.
PN WO200272771-A2.
PD 19-SEP-2002.
PA (CURA-) CURAGEN CORP.
Query Match
Best Local Similarity 8.7%; Score 147.5; DB 5; Length 2845;
21.8%; Pred. No. 0.01;
RESULT 1169
ID ABG22401 standard; protein; 361 AA.
DE Novel human diagnostic protein #22392.
PN WO200175067-A2.
PD 11-OCT-2001.
PA (HYSE-) HYSEQ INC.
Query Match
Best Local Similarity 8.7%; Score 146.5; DB 4; Length 361;
24.5%; Pred. No. 0.00078;
RESULT 1170
ID AAB85862 standard; protein; 373 AA.
DE Murine adipocytes-derived protein.
PN WO200166720-A1.
PD 13-SEP-2001.
PA (KITA/) KITAMURA T.
PA (TSUR/) TSURUGA H.
Query Match
Best Local Similarity 8.7%; Score 146.5; DB 4; Length 373;
25.9%; Pred. No. 0.00081;
RESULT 1171
ID AAB48126 standard; protein; 373 AA.
DE Mouse A236 polypeptide.
PN WO200069885-A2.
PD 23-NOV-2000.
PA (MILL-) MILLENNIUM PHARM INC.
Query Match
Best Local Similarity 8.6%; Score 145.5; DB 4; Length 373;
25.9%; Pred. No. 0.00099;
RESULT 1172
ID AAE26449 standard; protein; 373 AA.
DE Mouse A236 protein.
PN US2002055139-A1.
PD 09-MAY-2002.

PA (HOLT/) HOLTZMAN D A.
PA (SHAR/) SHARP J D.
PA (LEIB/) LEIBY K R.
PA (BOSS/) BOSSONE S.
PA (PANY/) PAN Y.
PA (BAEN/) BARNES T M.
PA (FRAS/) FRASER C C.
PA (WRIG/) WRIGHTON N.
PA (MYER/) MYERS P S.
PA (KING/) KINGSBURY G.

Query Match 8.6%; Score 145.5; DB 5; Length 373;
Best Local Similarity 25.9%; Pred. No. 0.00099;

RESULT 1173

ID ABB66424 standard; protein; 2016 AA.

DE Drosophila melanogaster polypeptide SEQ ID NO 26064.

PN WO200171042-A2.

PD 27-SEP-2001.

PA (PEKE) PE CORP NY.

Query Match 8.6%; Score 145; DB 4; Length 2016;
Best Local Similarity 25.6%; Pred. No. 0.011;

RESULT 1174

ID AAB48149 standard; protein; 373 AA.

DE Mouse A236 variant 2 polypeptide.

PN WO200069885-A2.

PD 23-NOV-2000.

PA (MILL-) MILLENNIUM PHARM INC.

Query Match 8.6%; Score 144.5; DB 4; Length 373;
Best Local Similarity 26.1%; Pred. No. 0.0012;

RESULT 1175

ID AAB48150 standard; protein; 373 AA.

DE Mouse A236 variant 3 polypeptide.

PN WO200069885-A2.

PD 23-NOV-2000.

PA (MILL-) MILLENNIUM PHARM INC.

Query Match 8.6%; Score 144.5; DB 4; Length 373;
Best Local Similarity 25.6%; Pred. No. 0.0012;

RESULT 1176

ID AAB48148 standard; protein; 373 AA.

DE Mouse A236 variant 1 polypeptide.

PN WO200069885-A2.

PD 23-NOV-2000.

PA (MILL-) MILLENNIUM PHARM INC.

Query Match 8.6%; Score 144.5; DB 4; Length 373;
Best Local Similarity 26.1%; Pred. No. 0.0012;

RESULT 1177

ID ADG63213 standard; protein; 367 AA.

DE Human neurotrophin protein +69bp isoform.

PN WO2003002765-A2.

PD 09-JAN-2003.

PA (IMCR) IMPERIAL CANCER RES TECHNOLOGY LTD.

Query Match 8.5%; Score 144; DB 7; Length 367;
Best Local Similarity 26.6%; Pred. No. 0.0013;

RESULT 1178

ID ABB63044 standard; protein; 467 AA.

DE Drosophila melanogaster polypeptide SEQ ID NO 15924.

PN WO200171042-A2.

PD 27-SEP-2001.

PA (PEKE) PE CORP NY.

Query Match 8.5%; Score 143.5; DB 4; Length 467;
Best Local Similarity 22.1%; Pred. No. 0.002;

RESULT 1179

ID ADE37401 standard; protein; 396 AA.

DE Murine nectin-like protein 1.

PN WO2003064992-A2.

PD 07-AUG-2003.

PA (PLAN-) PLANET BIOTECHNOLOGY INC.

PA (LARR-) LARRICK J W.

PA (WYCO-) WYCOFF K L.

Query Match 8.4%; Score 142.5; DB 7; Length 396;
Best Local Similarity 19.6%; Pred. No. 0.0019;

RESULT 1180

ID AAY69287 standard; protein; 398 AA.

DE Amino acid sequence of long extracellular form of murine B7-1 (CD80).

PN WO200008057-A2.

PD 17-FEB-2000.

PA (IMVX) IMMUNEX CORP.

Query Match 8.4%; Score 142.5; DB 3; Length 398;
Best Local Similarity 19.6%; Pred. No. 0.0019;

RESULT 1181

ID AAE00868 standard; protein; 404 AA.

DE Mouse brain immunoglobulin superfamily receptor (Bigr) protein.

PN WO200129083-A1.

PD 26-APR-2001.

PA (TEXA-) TEXAS BIOTECHNOLOGY CORP.

Query Match 8.4%; Score 142.5; DB 4; Length 404;
Best Local Similarity 19.6%; Pred. No. 0.002;

RESULT 1182

ID ABB68257 standard; protein; 1395 AA.

DE Drosophila melanogaster polypeptide SEQ ID NO 31563.

PN WO200171042-A2.

PD 27-SEP-2001.

PA (PEKE) PE CORP NY.

Query Match 8.4%; Score 142.5; DB 4; Length 1395;
Best Local Similarity 21.1%; Pred. No. 0.011;

RESULT 1183

ID AD808000 standard; protein; 376 AA.

DE Novel protein (useful for identifying genetic disorders) #155.

PN WO2003054152-A2.

PD 03-JUL-2003.

PA (HYSE-) HYSEQ INC.

Query Match 8.4%; Score 141.5; DB 7; Length 376;
Best Local Similarity 25.2%; Pred. No. 0.0022;

RESULT 1184

ID AAY13563 standard; protein; 1395 AA.

DE Drosophila Robo 1 polypeptide.

PN WO9925833-A1.

PD 27-MAY-1999.

PA (REGC) UNIV CALIFORNIA.

Query Match 8.4%; Score 141.5; DB 2; Length 1395;
Best Local Similarity 21.3%; Pred. No. 0.013;

RESULT 1185

ID AAY08401 standard; protein; 1395 AA.

DE Drosophila sp. Robo1 protein.

PN WO9920764-A1.

PD 29-APR-1999.

PA (REGC) UNIV CALIFORNIA.

Query Match 8.4%; Score 141.5; DB 2; Length 1395;
Best Local Similarity 21.3%; Pred. No. 0.013;

RESULT 1186

ID AD885335 standard; protein; 1395 AA.

DE Fruitfly nerve cell growth modulator SLIT-1-associated sequence #1.

PN US2003170727-A1.

PD 11-SEP-2003.

PA (GOOD/) GOODMAN C S.

PA (KIDD/) KIDD T.

PA (BROS/) BROSE K.

PA (TESS/) TESSIER-LAVIGNE M.

Query Match 8.4%; Score 141.5; DB 7; Length 1395;
Best Local Similarity 21.3%; Pred. No. 0.013;

RESULT 1187

ID AB084469 standard; protein; 3475 AA.

DE Human cancer-associated protein HP13-036.1.

PN WO2004074320-A2.

PD 02-SEP-2004.

PA (SAGR-) SAGRES DISCOVERY INC.

Query Match 8.4%; Score 141.5; DB 8; Length 3475;
Best Local Similarity 27.5%; Pred. No. 0.045;

RESULT 1188

ID ABB71502 standard; protein; 1052 AA.

DE Drosophila melanogaster polypeptide SEQ ID NO 41298.

PN WO200171042-A2.

PD 27-SEP-2001.

PA (PEKE) PE CORP NY.

Query Match 8.4%; Score 141; DB 4; Length 1052;
Best Local Similarity 25.1%; Pred. No. 0.0098;

RESULT 1189

ID AAW69698 standard; protein; 352 AA.
DE Mouse coxsackievirus and Ad2 and Ad5 receptor MCAR protein.
FN WO9833819-A1.
PD 06-AUG-1998.
PA (UUNY) UNIV NEW YORK STATE.
Query Match 8.3%; Score 140.5; DB 2; Length 352;
Best Local Similarity 25.4%; Pred. No. 0.0024;
RESULT 1190
ID AAW57213 standard; protein; 376 AA.
DE Mouse coxsackievirus and adenovirus receptor.
FN WO9811221-A2.
PD 19-MAR-1998.
PA (DAND) DANA FARBER CANCER INST INC.
Query Match 8.3%; Score 140.5; DB 2; Length 376;
Best Local Similarity 25.4%; Pred. No. 0.0027;
RESULT 1191
ID ADP56685 standard; protein; 265 AA.
DE Human junction adhesion molecule 3 splice variant 2 (huJAM3sv2) protein.
FN WO2004053058-A2.
PD 24-JUN-2004.
PA (ELIL) LILLY & CO ELI.
Query Match 8.3%; Score 139.5; DB 8; Length 265;
Best Local Similarity 24.0%; Pred. No. 0.002;
RESULT 1192
ID AAY96294 standard; protein; 310 AA.
DE Human IGFAM-6 immunoglobulin.
FN WO200029583-A2.
PD 25-MAY-2000.
PA (INCY-) INCYTE PHARM INC.
Query Match 8.3%; Score 139.5; DB 3; Length 310;
Best Local Similarity 22.0%; Pred. No. 0.0025;
RESULT 1193
ID ADP56683 standard; protein; 310 AA.
DE Human junction adhesion molecule 3 (huJAM3) full-length protein.
FN WO2004053058-A2.
PD 24-JUN-2004.
PA (ELIL) LILLY & CO ELI.
Query Match 8.3%; Score 139.5; DB 8; Length 310;
Best Local Similarity 22.0%; Pred. No. 0.0025;
RESULT 1194
ID AAY96735 standard; protein; 310 AA.
DE PRO1868, an A33 antigen homologue.
FN WO200036102-A2.
PD 22-JUN-2000.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 3; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1195
ID AAB33457 standard; protein; 310 AA.
DE Human PRO1868 protein UNQ859 SEQ ID NO:193.
FN WO200053758-A2.
PD 14-SEP-2000.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 3; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1196
ID AAB27276 standard; protein; 310 AA.
DE Human confluency regulated adhesion molecule 1 #2.
FN WO200053749-A2.
PD 14-SEP-2000.
PA (RMFD-) RMF DICTAGENE SA.
Query Match 8.2%; Score 138.5; DB 3; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1197
ID AAB80272 standard; protein; 310 AA.
DE Human PRO1868 protein.
FN WO200104311-A1.
PD 18-JAN-2001.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 4; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1198
ID AAW93905 standard; protein; 310 AA.

DE Human polypeptide, SEQ ID NO: 4051.
FN EP1130094-A2.
PD 05-SEP-2001.
PA (HELI-) HELIX RES INST.
Query Match 8.2%; Score 138.5; DB 4; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1199
ID AAM93323 standard; protein; 310 AA.
DE Human polypeptide, SEQ ID NO: 2845.
FN EP1130094-A2.
PD 05-SEP-2001.
PA (HELI-) HELIX RES INST.
Query Match 8.2%; Score 138.5; DB 4; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1200
ID AAU12440 standard; protein; 310 AA.
DE Human PRO1868 polypeptide sequence.
FN WO200140466-A2.
PD 07-JUN-2001.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 4; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1201
ID AAB80383 standard; protein; 310 AA.
DE Secreted protein encoded by gene #13.
FN WO200107459-A1.
PD 01-FEB-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 4; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1202
ID AAB80408 standard; protein; 310 AA.
DE Secreted protein encoded by gene #38.
FN WO200107459-A1.
PD 01-FEB-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 4; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1203
ID AAB80409 standard; protein; 310 AA.
DE Secreted protein encoded by gene #39.
FN WO200107459-A1.
PD 01-FEB-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 4; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1204
ID ABG92709 standard; protein; 310 AA.
DE Human secreted protein PRO1868.
FN US2002098506-A1.
PD 25-JUL-2002.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 5; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1205
ID ABG91361 standard; protein; 310 AA.
DE Novel human secreted protein #7.
FN US2002098505-A1.
PD 25-JUL-2002.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 5; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1206
ID ABB84947 standard; protein; 310 AA.
DE Human PRO1868 protein sequence SEQ ID NO:262.
FN WO200200690-A2.
PD 03-JAN-2002.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 5; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1207
ID ABG65297 standard; protein; 310 AA.
DE Human albumin fusion protein #1972.

PN WO200177137-A1.
 PD 18-OCT-2001.
 PA (HUMA-) HUMAN GENOME SCI INC.
 Query Match 8.2%; Score 138.5; DB 5; Length 310;
 Best Local Similarity 21.8%; Pred. No. 0.0031;
 RESULT 1208
 ID ABG65296 standard; protein; 310 AA.
 DE Human albumin fusion protein #1971.
 PN WO200177137-A1.
 PD 18-OCT-2001.
 PA (HUMA-) HUMAN GENOME SCI INC.
 Query Match 8.2%; Score 138.5; DB 5; Length 310;
 Best Local Similarity 21.8%; Pred. No. 0.0031;
 RESULT 1209
 ID ABG65298 standard; protein; 310 AA.
 DE Human albumin fusion protein #1973.
 PN WO200177137-A1.
 PD 18-OCT-2001.
 PA (HUMA-) HUMAN GENOME SCI INC.
 Query Match 8.2%; Score 138.5; DB 5; Length 310;
 Best Local Similarity 21.8%; Pred. No. 0.0031;
 RESULT 1210
 ID ABG31401 standard; protein; 310 AA.
 DE Human PRO1868 polypeptide.
 PN US2002098507-A1.
 PD 25-JUL-2002.
 PA (GETH) GENENTECH INC.
 Query Match 8.2%; Score 138.5; DB 5; Length 310;
 Best Local Similarity 21.8%; Pred. No. 0.0031;
 RESULT 1211
 ID ABB95553 standard; protein; 310 AA.
 DE Human angiogenesis related protein PRO1868 SEQ ID NO: 262.
 PN WO200208284-A2.
 PD 31-JAN-2002.
 PA (GETH) GENENTECH INC.
 Query Match 8.2%; Score 138.5; DB 5; Length 310;
 Best Local Similarity 21.8%; Pred. No. 0.0031;
 RESULT 1212
 ID ABU71650 standard; protein; 310 AA.
 DE Human PRO polypeptide #61.
 PN US2002146709-A1.
 PD 10-OCT-2002.
 PA (GETH) GENENTECH INC.
 Query Match 8.2%; Score 138.5; DB 6; Length 310;
 Best Local Similarity 21.8%; Pred. No. 0.0031;
 RESULT 1213
 ID ABU72377 standard; protein; 310 AA.
 DE Novel human secreted and transmembrane protein PRO1868.
 PN US2002182618-A1.
 PD 05-DEC-2002.
 PA (GETH) GENENTECH INC.
 Query Match 8.2%; Score 138.5; DB 6; Length 310;
 Best Local Similarity 21.8%; Pred. No. 0.0031;
 RESULT 1214
 ID ABUS0867 standard; protein; 310 AA.
 DE Human secreted and transmembrane polypeptide PRO1868.
 PN US2002192868-A1.
 PD 19-DEC-2002.

PA (GETH) GENENTECH INC.
 Query Match 8.2%; Score 138.5; DB 6; Length 310;
 Best Local Similarity 21.8%; Pred. No. 0.0031;
 RESULT 1215
 ID ABO17884 standard; protein; 310 AA.
 DE Novel human secreted and transmembrane protein PRO1868.
 PN US2003032156-A1.
 PD 13-FEB-2003.
 PA (GETH) GENENTECH INC.
 Query Match 8.2%; Score 138.5; DB 6; Length 310;
 Best Local Similarity 21.8%; Pred. No. 0.0031;
 RESULT 1216
 ID ABU71505 standard; protein; 310 AA.
 DE Human PRO polypeptide #61.
 PN US2002192659-A1.
 PD 19-DEC-2002.
 PA (GETH) GENENTECH INC.
 Query Match 8.2%; Score 138.5; DB 6; Length 310;
 Best Local Similarity 21.8%; Pred. No. 0.0031;
 RESULT 1217
 ID ADA57610 standard; protein; 310 AA.
 DE Human secreted protein #592.
 PN WO2002102994-A2.
 PD 27-DEC-2002.
 PA (HUMA-) HUMAN GENOME SCI INC.
 Query Match 8.2%; Score 138.5; DB 6; Length 310;
 Best Local Similarity 21.8%; Pred. No. 0.0031;
 RESULT 1218
 ID ADA57611 standard; protein; 310 AA.
 DE Human secreted protein #592.
 PN WO2002102994-A2.
 PD 27-DEC-2002.
 PA (HUMA-) HUMAN GENOME SCI INC.
 Query Match 8.2%; Score 138.5; DB 6; Length 310;
 Best Local Similarity 21.8%; Pred. No. 0.0031;
 RESULT 1219
 ID ADA57309 standard; protein; 310 AA.
 DE Human secreted protein #592.
 PN WO2002102994-A2.
 PD 27-DEC-2002.
 PA (HUMA-) HUMAN GENOME SCI INC.
 Query Match 8.2%; Score 138.5; DB 6; Length 310;
 Best Local Similarity 21.8%; Pred. No. 0.0031;
 RESULT 1220
 ID ABP71277 standard; protein; 310 AA.
 DE Human junctional adhesion molecule 3 (JAM3).
 PN WO2003006673-A2.
 PD 23-JAN-2003.
 PA (TEXA-) TEXAS BIOTECHNOLOGY CORP.
 Query Match 8.2%; Score 138.5; DB 6; Length 310;
 Best Local Similarity 21.8%; Pred. No. 0.0031;
 RESULT 1221
 ID ASU81138 standard; protein; 310 AA.
 DE Human PRO polypeptide #269.
 PN US2003004311-A1.
 PD 02-JAN-2003.
 PA (GETH) GENENTECH INC.
 Query Match 8.2%; Score 138.5; DB 6; Length 310;
 Best Local Similarity 21.8%; Pred. No. 0.0031;
 RESULT 1222
 ID ABU71951 standard; protein; 310 AA.
 DE Human secreted/transmembrane protein PRO1868.
 PN US2003003530-A1.
 PD 02-JAN-2003.
 PA (GETH) GENENTECH INC.
 Query Match 8.2%; Score 138.5; DB 6; Length 310;
 Best Local Similarity 21.8%; Pred. No. 0.0031;
 RESULT 1223
 ID ABO01834 standard; protein; 310 AA.
 DE Novel human secreted and transmembrane protein PRO1868.
 PN US2002197671-A1.
 PD 26-DEC-2002.
 PA (GETH) GENENTECH INC.

Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1224
ID ABU66838 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003036180-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1225
ID ABU54407 standard; protein; 310 AA.
DE Human secreted/transmembrane protein PRO1868.
PN US2002132240-A1.
PD 19-SEP-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1226
ID AB047422 standard; protein; 310 AA.
DE Human secreted/transmembrane polypeptide PRO1868.
PN US2003044839-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1227
ID ABG73314 standard; protein; 310 AA.
DE Human PRO1868 polypeptide.
PN US2002164646-A1.
PD 07-NOV-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1228
ID ABU59919 standard; protein; 310 AA.
DE Novel secreted and transmembrane protein PRO1868.
PN US2003017563-A1.
PD 23-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1229
ID AB025109 standard; protein; 310 AA.
DE Human secreted/transmembrane protein (PRO) #269.
PN US2003036179-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1230
ID ABU64559 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #63.
PN US2002160374-A1.
PD 31-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1231
ID ABU67405 standard; protein; 310 AA.
DE Human secreted protein PRO1868.
PN US2003023054-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1232
ID AB014925 standard; protein; 310 AA.
DE Human secreted / transmembrane polypeptide PRO1868.
PN US2003036060-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;

Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1233
ID ABU60813 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #7.
PN US2002160392-A1.
PD 31-OCT-2002.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1234
ID ABU67114 standard; protein; 310 AA.
DE Human secreted/transmembrane, PRO, protein SEQ ID 538.
PN US2003032155-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1235
ID ABU81236 standard; protein; 310 AA.
DE Human PRO1917polypeptide.
PN US2003032060-A1.
PD 13-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1236
ID ABU69682 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868+H30.
PN US2003017463-A1.
PD 23-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1237
ID ABQ14864 standard; protein; 310 AA.
DE Human secreted / transmembrane polypeptide PRO1868.
PN US2003027143-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1238
ID ADA46057 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003022328-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1239
ID ADA76488 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003073212-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1240
ID ADB29627 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003092002-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1241
ID ADA19138 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003054517-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1242
ID ADB29627 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003092002-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;
RESULT 1243
ID AB014925 standard; protein; 310 AA.
DE Human secreted / transmembrane polypeptide PRO1868.
PN US2003036060-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 6; Length 310;

RESULT 1242
ID ADA61761 standard; protein; 310 AA.
DE Homo sapiens.
PN US2003049816-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1243
ID ADB19546 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003068796-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1244
ID ADB28087 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082704-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1245
ID ADA86566 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082711-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1246
ID ADB16130 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003087350-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1247
ID ADA47916 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003073215-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1248
ID ADA18484 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003039971-A1.
PD 27-FEB-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1249
ID ABO32816 standard; protein; 310 AA.
DE Human secreted/transmembrane protein PRO1868.
PN US2003045693-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1250
ID ADA67711 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003068795-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1251

ID ADB30718 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003068794-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1252
ID ADA86014 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082693-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1253
ID ADA97226 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082705-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1254
ID ADA79530 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082763-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1255
ID ADA87669 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087345-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1256
ID ADB16871 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003087349-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1257
ID ABO34876 standard; protein; 310 AA.
DE Human PRO polypeptide #61.
PN US2003044793-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1258
ID ADA16459 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003049621-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1259
ID ADA91963 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082694-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 6; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1260
ID ADB15026 standard; protein; 310 AA.

DE Human PRO polypeptide #269.
PN US2003087351-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1261
ID ADB18987 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003073211-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1262
ID ADA94202 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003077722-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1263
ID ADB20098 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082691-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1264
ID ADB13410 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082710-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1265
ID ABO43417 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US200304945-A1.
PD 06-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1266
ID ADA74664 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003068798-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1267
ID ADA42604 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003054401-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1268
ID ADB24897 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US200307713-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1269
ID ADA82421 standard; protein; 310 AA.
DE Human PRO polypeptide #269.

PN US2003082701-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1270
ID ADA75384 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003073216-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1271
ID ADA85462 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082695-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1272
ID ADA84910 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082708-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1273
ID ABO17554 standard; protein; 310 AA.
DE Human PRO polypeptide #61.
PN US2003064367-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1274
ID ADB30166 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003073214-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1275
ID ADA80694 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082761-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1276
ID ADA75936 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082703-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1277
ID ADA47161 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003073210-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1278
ID ADB25457 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US2003077715-A1.

PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1279
ID ADA93633 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US200307721-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1280
ID ADB26983 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003092147-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1281
ID ADB31270 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003096386-A1.
PD 22-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1282
ID ABU62957 standard; protein; 310 AA.
DE Human PRO1868 protein.
PN US2003054447-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1283
ID ADA61198 standard; protein; 310 AA.
DE Homo sapiens.
PN US2003049817-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1284
ID ADB24345 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US200307714-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1285
ID ADA96674 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082690-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1286
ID ADA81246 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082702-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1287
ID ADA96122 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082759-A1.
PD 01-MAY-2003.

PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1288
ID ADB26431 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082760-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 6; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1289
ID ADB21916 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082765-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1290
ID ADA77695 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003068797-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1291
ID ADB18435 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US200307710-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1292
ID ADA87118 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082709-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1293
ID ADA16883 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003039969-A1.
PD 27-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1294
ID ADA13312 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003049622-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1295
ID ADA42180 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003082540-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1296
ID ADA88221 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082700-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.

Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1297
ID ADA46609 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003054516-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1298
ID ADA17527 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003017498-A1.
PD 23-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1299
ID ADA43030 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003054351-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1300
ID ADB28639 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082699-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1301
ID ADB29191 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082706-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1302
ID ABO01894 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003027256-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1303
ID ADA77143 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003059909-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1304
ID ADA88773 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003073213-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1305
ID ADA97778 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082686-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1306
ID ADB27535 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003022239-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1307
ID ADB22468 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087344-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1308
ID ABO17615 standard; protein; 310 AA.
DE Human PRO polypeptide #61.
PN US2003064923-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1309
ID ADA67159 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003068793-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1310
ID ADB23020 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003077711-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1311
ID ADB23793 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US2003077712-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1312
ID ADA92515 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082712-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1313
ID ADB15578 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003087352-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1314
ID ADB38830 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082766-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1315
ID ADB38278 standard; protein; 310 AA.

Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1306
ID ADB27535 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003022239-A1.
PD 30-JAN-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1307
ID ADB22468 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087344-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1308
ID ABO17615 standard; protein; 310 AA.
DE Human PRO polypeptide #61.
PN US2003064923-A1.
PD 03-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1309
ID ADA67159 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003068793-A1.
PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1310
ID ADB23020 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003077711-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1311
ID ADB23793 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US2003077712-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1312
ID ADA92515 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082712-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1313
ID ADB15578 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003087352-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1314
ID ADB38830 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082766-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 8.2%; Score 138.5; DB 7; Length 310;
RESULT 1315
ID ADB38278 standard; protein; 310 AA.

DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087347-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1316
ID ADB66750 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082689-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1317
ID ADB89830 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082698-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1318
ID ADB90562 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082762-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1319
ID ADB77948 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003077654-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1320
ID ADB39663 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082764-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1321
ID ADB75084 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003082542-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1322
ID ADB47286 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082687-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1323
ID ADB86893 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003082697-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1324
ID ADB77498 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.

PN US2003082696-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1325
ID ADB34655 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US2003077717-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1326
ID ADB35759 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US2003077719-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1327
ID ADB34103 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US2003077716-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1328
ID ADB35207 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US2003077718-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1329
ID ADB36311 standard; protein; 310 AA.
DE Human PRO polypeptide SEQ ID NO 538.
PN US2003077720-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1330
ID ADB46706 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003082692-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1331
ID ADC28731 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003059772-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1332
ID ADC39931 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003059828-A1.
PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1333
ID ADC40445 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003059829-A1.

PD 27-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1334
ID ADC19269 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003036061-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1335
ID ADC34569 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003036094-A1.
PD 20-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1336
ID ADC29624 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003049676-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1337
ID ADC29155 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003049677-A1.
PD 13-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1338
ID ADC41040 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003054400-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1339
ID ADC19697 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003054441-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1340
ID ADC34145 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003073077-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1341
ID ADC13215 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003073079-A1.
PD 17-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1342
ID AAE38826 standard; protein; 310 AA.
DE Human PRO1868 protein.
PN US200307657-A1.
PD 24-APR-2003.

PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1343
ID ADC50579 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003092106-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1344
ID ADC72126 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003092107-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1345
ID ADC60105 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003092105-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1346
ID ADC53112 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein Seq ID538.
PN US2003087365-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1347
ID ADC57466 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein Seq ID538.
PN US2003087366-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1348
ID ADC60657 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087367-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1349
ID ADC51132 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087361-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1350
ID ADC65659 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003087362-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1351
ID ADC54757 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein Seq ID538.
PN US2003087363-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.

Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1352
ID ADC53718 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein Seq ID538.
PN US2003087364-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1353
ID ADC59241 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein Seq ID538.
PN US2003087359-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1354
ID ADC56119 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein Seq ID538.
PN US2003087360-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1355
ID ADC58689 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein Seq ID538.
PN US2003087346-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1356
ID ADC2667 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003082541-A1.
PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1357
ID ADC74383 standard; protein; 310 AA.
DE Human secreted protein - SEQ ID 1016.
PN WO2003038063-A2.
PD 08-MAY-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1358
ID ADC74606 standard; protein; 310 AA.
DE Human secreted protein - SEQ ID 1239.
PN WO2003038083-A2.
PD 08-MAY-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1359
ID ADC74607 standard; protein; 310 AA.
DE Human secreted protein - SEQ ID 1240.
PN WO2003038063-A2.
PD 08-MAY-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1360
ID ADD03363 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003092104-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;

Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1361
ID ADC90355 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087348-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1362
ID ADC69774 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194770-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1363
ID ADC48663 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194773-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1364
ID ADD10192 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194776-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1365
ID ADD04767 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087354-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1366
ID ADC80723 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003092103-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1367
ID ADD11230 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194774-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1368
ID ADD10551 standard; protein; 310 AA.
DE Human secreted/transmembrane PRO polypeptide #131.
PN US2003105011-A1.
PD 05-JUN-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1369
ID ADC48111 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194771-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;

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RESULT 1370
ID ADD05222 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003104469-A1.
PD 05-JUN-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1371
ID ADC80171 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087358-A1.
PD 08-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1372
ID ADD11511 standard; protein; 310 AA.
DE Human secreted/transmembrane PRO polypeptide #131.
PN US2003105013-A1.
PD 05-JUN-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1373
ID ADD09640 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194775-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1374
ID ADD04228 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003104381-A1.
PD 05-JUN-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1375
ID ADD03804 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003108983-A1.
PD 12-JUN-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1376
ID ADD41353 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003203438-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1377
ID ADD52492 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194769-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1378
ID ADD53232 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194792-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1379
ID ADD92672 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003203432-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1380
ID ADD37304 standard; protein; 310 AA.
DE Human secreted/transmembrane PRO polypeptide #131.
PN US2003105012-A1.
PD 05-JUN-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1381
ID ADD51940 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194779-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1382
ID ADD02739 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003203431-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1383
ID ADD38106 standard; protein; 310 AA.
DE Human secreted protein #289.
PN WO200290526-A2.
PD 14-NOV-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1384
ID ADD38009 standard; protein; 310 AA.
DE Human secreted protein #192.
PN WO200290526-A2.
PD 14-NOV-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1385
ID ADD38105 standard; protein; 310 AA.
DE Human secreted protein #288.
PN WO200290526-A2.
PD 14-NOV-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1386
ID ADD02173 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003203430-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1387
ID ADD54355 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003203432-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1388
ID ADD92672 standard; protein; 310 AA.
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DE Human PRO polypeptide #269.
PN US2003199030-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1389
ID ADD91568 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199055-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1390
ID ADE04182 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199057-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1391
ID ADE32479 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003194765-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1392
ID ADE22411 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199056-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1393
ID ADD79635 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003203428-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1394
ID ADE42171 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194772-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1395
ID ADE17988 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199023-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1396
ID ADD92120 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199053-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1397
ID ADE33583 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.

PN US2003194767-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1398
ID ADE34135 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003194791-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1399
ID ADD80187 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207417-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1400
ID ADD93224 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194768-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1401
ID ADE19644 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199025-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1402
ID ADE35056 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003077583-A1.
PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1403
ID ADE19092 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199026-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1404
ID ADE43288 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199033-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1405
ID ADD96077 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199059-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1406
ID ADE22963 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199064-A1.

PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1407
ID ADD79081 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003203439-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1408
ID ADE33031 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003194766-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1409
ID ADE42723 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199032-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1410
ID ADD80739 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207418-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1411
ID ADD89767 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199028-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1412
ID ADE41051 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199031-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1413
ID ADE04850 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199034-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1414
ID ADE92979 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194777-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1415
ID ADG21688 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207355-A1.
PD 06-NOV-2003.

PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1416
ID ADG23329 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207384-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1417
ID ADF97664 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207370-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1418
ID ADG80728 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207373-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1419
ID ADG80176 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207372-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1420
ID ADH62558 standard; protein; 310 AA.
DE Human PRO1868 protein.
PN US2003171568-A1.
PD 11-SEP-2003.
PA (ASHK/) ASHKENAZI A.
PA (FONG/) FONG S.
PA (GODD/) GODDARD A.
PA (GURN/) GURNEY A L.
PA (NAPI/) RAPIER M A.
PA (TUNA/) TOMAS D.
PA (WOOD/) WOOD W I.
Query Match Similarity 21.8%; Score 138.5; DB 7; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1421
ID ADH59539 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003039972-A1.
PD 27-FEB-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1422
ID ADH55468 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207381-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1423
ID ADH56020 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207379-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1424
ID ADG21688 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207355-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Similarity 21.8%; Pred. No. 0.0031;
Best Local Similarity 21.8%; Pred. No. 0.0031;

RESULT 1424
ID AD138318 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003054352-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1425
ID AD164239 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207385-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1426
ID AD165188 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207386-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1427
ID AD163687 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207387-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1428
ID ADH82101 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207388-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1429
ID ADH1549 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207377-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1430
ID ADJ58518 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003170864-A1.
PD 11-SEP-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1431
ID ADJ26586 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003054349-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1432
ID ADM82718 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087355-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1433

ID ADM16117 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087353-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1434
ID ADM16746 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087385-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1435
ID ADM15565 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087356-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1436
ID ADM15013 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003087357-A1.
PD 08-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 7; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1437
ID ADC81275 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003092115-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 8; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1438
ID ADE79501 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003135025-A1.
PD 17-JUL-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 8; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1439
ID ADD76723 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003100087-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 8; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1440
ID ADD88087 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003092113-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 8; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1441
ID ADD86491 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003203440-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC. 8.2%; Score 138.5; DB 8; Length 310;
Query Match Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1442
ID ADE79925 standard; protein; 310 AA.

DE Human secreted/transmembrane protein, #65.
PN US2003130489-A1.
PD 10-JUL-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1443
ID ADE75939 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003211571-A1.
PD 13-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1444
ID ADE73601 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003129592-A1.
PD 10-JUL-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1445
ID ADE41512 standard; protein; 310 AA.
DE Human secreted/transmembrane PRO polypeptide #131.
PN US2003100497-A1.
PD 29-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1446
ID ADE23515 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003092108-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1447
ID ADE24067 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003092110-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1448
ID ADE24710 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003092111-A1.
PD 15-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1449
ID ADD87535 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003203439-A1.
PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1450
ID ADE89401 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199062-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1451
ID ADE74136 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.

PN US2003148370-A1.
PD 07-AUG-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1452
ID ADE18540 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003194794-A1.
PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1453
ID ADE88849 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199054-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1454
ID ADE99690 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003211576-A1.
PD 13-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1455
ID ADE94869 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199027-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1456
ID ADE91280 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199061-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1457
ID ADE95421 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199052-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1458
ID ADE93531 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199060-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1459
ID ADF35112 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199029-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1460
ID ADE98809 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003211569-A1.

PD 13-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1461
ID ADE92427 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003199051-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1462
ID ADE90728 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003199063-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1463
ID ADE91875 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003199058-A1.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1464
ID ADE99236 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003211568-A1.
PD 13-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1465
ID ADG40706 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003225253-A1.
PD 04-DEC-2003.
PA (DESN/) DESNOYERS L.
PA (GODD/) GODDARD A.
PA (GODO/) GODOWSKI P J.
PA (GURN/) GURNEY A L.
PA (MATH/) MATHER J P.
PA (WILL/) WILLIAMS P M.
PA (WOOD/) WOOD W I.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1466
ID ADF74100 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003180312-A1.
PD 25-SEP-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1467
ID ADG2454 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207352-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1468
ID ADG22240 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207360-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1469
ID ADG20310 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207376-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1470
ID ADF98216 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207422-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1471
ID ADG24433 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207426-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1472
ID ADF98787 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003208055-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1473
ID ADG03618 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207351-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1474
ID ADF99339 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207353-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1475
ID ADG16924 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207359-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1476
ID ADG05383 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207375-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1477
ID ADG19650 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207425-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;

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RESULT 1478
ID ADF73676 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003166051-A1.
PD 04-SEP-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1479
ID ADG13487 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207357-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1480
ID ADG08544 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207424-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1481
ID ADG15714 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003219885-A1.
PD 27-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1482
ID ADF97112 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207371-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1483
ID ADG06297 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207374-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1484
ID ADG23881 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207389-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1485
ID ADG04170 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207423-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1486
ID ADG25071 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207427-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1487
ID ADG55967 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207350-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1488
ID ADG07920 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207356-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1489
ID ADG55415 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003194778-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1490
ID ADG61079 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207390-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1491
ID ADG62183 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207428-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1492
ID ADG92519 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003027145-A1.
PD 06-FEB-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1493
ID ADG82384 standard; protein; 310 AA.
DE Human PRO polypeptide #269.
PN US2003207358-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1494
ID ADG57623 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207362-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1495
ID ADG57071 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207364-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
```


DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207365-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1497
ID ADG58727 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207368-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1498
ID ADG71093 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207420-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1499
ID ADG92946 standard; protein; 310 AA.
DE Human secreted/transmembrane protein, #65.
PN US2003027146-A1.
PD 06-FEB-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;
RESULT 1500
ID ADG58175 standard; protein; 310 AA.
DE Novel human secreted and transmembrane protein PRO1868.
PN US2003207363-A1.
PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 8.2%; Score 138.5; DB 8; Length 310;
Best Local Similarity 21.8%; Pred. No. 0.0031;

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GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: August 26, 2005, 19:28:02 ; Search time 40 Seconds
(without alignments)
772.140 Million cell updates/sec

Title: US-10-767-374-2

Perfect score: 1688

Sequence: 1 MGILGLLLGLHGLTVDTYGR.....AYIMLCRKTSQOEHVYEAAAR 321

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 283416 seqs, 96216763 residues

Total number of hits satisfying chosen parameters: 283416

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-Processing: Minimum Match 0%

Maximum Match 100%

Listing first 1500 summaries

Database : PIR 79:*

1: pir1:*

2: pir2:*

3: pir3:*

4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	178.5	10.6	299	2 S56749	junctional adhesio
2	172	10.2	7962	2 I38346	elastic titin - hu
3	169	10.0	365	2 JC7780	coxsackie- and ade
4	143.5	8.5	1052	2 B49120	protein-tyrosine k
5	139.5	8.3	3707	2 S18252	heparan sulfate pr
6	138.5	8.2	1323	2 PN0568	connectin 3B - chi
7	138.5	8.2	4162	2 T42633	connectin/titin -
8	136	8.1	584	2 T08678	hypothetical prote
9	133.5	7.9	5175	2 T20992	hypothetical prote
10	133.5	7.9	5198	2 T43290	hemiscentin precurs
11	133	7.9	1106	1 PFHUGB	platelet-derived g
12	132.5	7.8	765	2 C42632	cell adhesion mole
13	132.5	7.8	812	2 B42632	cell adhesion mole
14	132.5	7.8	932	2 A42632	cell adhesion mole
15	132.5	7.8	1272	2 S26180	neurofascin - chic
16	132	7.8	1896	2 T08851	Down syndrome cell
17	131	7.8	4391	2 A38051	perlecan precursor
18	129.5	7.7	1040	2 A34695	axonal glycoprotei
19	129.5	7.7	3375	2 T19821	hypothetical prote
20	129	7.6	1427	2 I51669	tumor suppressor -
21	128.5	7.6	338	2 JC5519	50K glycoprotein p
22	127.5	7.6	1906	1 S68235	myosin-light-chain
23	125.5	7.4	1259	2 A43425	Bravo/Nr-CAM cell
24	125	7.4	351	2 JH0396	biliary glycoprote
25	124	7.3	853	1 IJBONC	neural cell adhesi
26	124	7.3	1344	2 T14316	rig-1 protein - mo
27	123.5	7.3	458	1 WMMSR1	biliary glycoprote
28	123.5	7.3	521	2 JC1508	biliary glycoprote
29	123.5	7.3	1091	1 IJCHNL	neural cell adhesi

30	122.5	7.3	761	1 IJHUNG	neural cell adhesi
31	122.5	7.3	1268	1 A39640	neural cell adhesi
32	120.5	7.1	458	2 JC1509	biliary glycoprote
33	120.5	7.1	538	2 JC2457	vascular cell adhe
34	120	7.1	1033	2 S19247	cell adhesion prot
35	120	7.1	1036	2 S22383	axonin 1 precursor
36	119.5	7.1	417	2 JH0394	biliary glycoprote
37	119.5	7.1	464	2 C30127	transmembrane carc
38	119.5	7.1	483	2 T17346	hypothetical prote
39	119.5	7.1	526	1 A32164	biliary glycoprote
40	119.5	7.1	725	1 IJMSNG	neural cell adhesi
41	119.5	7.1	725	2 JH0100	neural cell adhesi
42	119.5	7.1	1115	1 IJMSNL	neural cell adhesi
43	119	7.0	467	1 HLMSP3	poliovirus recepto
44	119	7.0	1091	2 A58532	glial cell membran
45	119	7.0	26926	1 I38344	titin, cardiac mus
46	118.5	7.0	333	2 A31923	amalgam protein pr
47	118.5	7.0	458	2 S68177	C-CAM2a protein is
48	118.5	7.0	458	2 S23969	cell-adhesion mole
49	118.5	7.0	519	2 A44783	ecto-ATPase precu
50	118.5	7.0	816	2 A49151	fibroblast growth
51	118.5	7.0	1011	2 T13669	neuromusculin - fr
52	118.5	7.0	1694	2 S50065	sialoadhesin - mou
53	118	7.0	547	1 S28904	intercellular adhe
54	118	7.0	1338	2 S09982	protein-tyrosine k
55	117.5	7.0	338	2 JC4776	limbic-system-asso
56	117.5	7.0	344	2 A27681	nonspecific cross-
57	117.5	7.0	1092	1 JN0635	neural cell adhesi
58	117.5	7.0	1239	1 A32579	neuroglian - fruit
59	117	6.9	739	2 J80675	vascular cell adhe
60	117	6.9	858	1 IJRTNC	neural cell adhesi
61	117	6.9	1447	2 A54100	tumor suppressor p
62	116.5	6.9	806	2 A35963	protein-tyrosine k
63	116.5	6.9	1040	2 A49356	transient axonal g
64	116.5	6.9	1051	2 A39712	kinase-like protei
65	116	6.9	344	2 I56551	neurotrophin rat
66	116	6.9	6658	2 T13931	projectin - fruit
67	115.5	6.8	292	2 S03421	T-cell receptor de
68	115.5	6.8	293	2 A40131	T-cell receptor de
69	115.5	6.8	309	2 I49522	gene B7-2 protein
70	115.5	6.8	725	2 J80099	neural cell adhesi
71	115.5	6.8	738	2 A40096	platelet-endotheli
72	115.5	6.8	1257	1 A41060	neural cell adhesi
73	115	6.8	1209	2 T42718	probable neural ce
74	114.5	6.8	478	2 I53960	PRR2 alpha - human
75	114.5	6.8	521	2 S34338	biliary glycoprote
76	114.5	6.8	538	2 I68093	PRR2 delta - human
77	114.5	6.8	1088	1 IJXLNL	neural cell adhesi
78	114.5	6.8	1173	2 T25893	hypothetical prote
79	114	6.8	544	2 JC5018	intercellular adhe
80	113.5	6.7	530	2 A53437	poliovirus recepto
81	113	6.7	1277	2 T30532	neural cell adhesi
82	112.5	6.7	321	2 JH0395	biliary glycoprote
83	112.5	6.7	345	2 JC4025	opioid-binding cel
84	112.5	6.7	1259	2 S36126	neural cell adhesi
85	112.5	6.7	2029	1 TDFFLK	protein-tyrosine-p
86	112	6.6	946	1 A47299	ror-related recept
87	112	6.6	976	2 T29583	hypothetical prote
88	112	6.6	1273	2 T42405	sax-3 protein - Ca
89	111	6.6	806	1 TVHUF3	fibroblast growth
90	111	6.6	6642	2 T29757	protein UNC-89 - C
91	110.5	6.5	349	2 A34815	carcinoembryonic a
92	110.5	6.5	518	2 JC4024	poliovirus recepto
93	110	6.5	499	2 S33766	platelet-derived g
94	110	6.5	917	2 I48950	telencephalin prec
95	110	6.5	1070	2 JC4593	protein-tyrosine k
96	110	6.5	1880	2 T18531	tractin - medicina
97	110	6.5	2295	2 C88369	protein unc-52 [im
98	109.5	6.5	338	2 JC1238	opioid-binding pro
99	109.5	6.5	345	2 JC1239	opioid-binding pro
100	109.5	6.5	526	2 S70587	butyrophilin precu
101	109.5	6.5	1336	2 I60598	Fit-1 tyrosine kin
102	109	6.5	547	2 B45808	B-lymphocyte antig

103	109	6.5	646	2	I38049	cell surface glyco	176	100.5	6.0	802	1	TVHUF4	fibroblast growth
104	108.5	6.4	345	2	S03199	opioid-binding pro	177	100.5	6.0	847	2	JH0371	B-cell adhesion pr
105	108.5	6.4	416	2	A54017	colon carcinoma-as	178	100	5.9	354	1	LKHU	proteoglycan link
106	108.5	6.4	1328	2	T23007	hypothetical prote	179	100	5.9	520	1	S44099	brain-derived neur
107	108	6.4	526	1	A37821	butyrophilin - bov	180	100	5.9	818	1	S44098	brain-derived neur
108	107	6.3	773	1	QRREG	secretory componen	181	100	5.9	1021	2	I39207	leukocyte surface
109	106.5	6.3	354	1	S42938	proteoglycan link	182	99.5	5.9	459	2	A46254	CD4 precursor - ra
110	106.5	6.3	1227	2	T23004	hypothetical prote	183	99.5	5.9	626	1	A61084	myelin-associated
111	106.5	6.3	1232	2	T32021	neural cell adhesi	184	99.5	5.9	790	2	A39627	protein-tyrosine k
112	106	6.3	319	2	T32541	unc-5 protein - Ca	185	99.5	5.9	1028	2	I58164	BIG-1 protein - ra
113	106	6.3	947	1	B44294	unc-5 protein, lon	186	99.5	5.9	1379	2	JC4954	vascular endotheli
114	106	6.3	1040	2	A57638	receptor tyrosine	187	99.5	5.9	1666	2	A48594	skelemin - mouse
115	106	6.3	1098	1	PFMSRB	platelet-derived g	188	99	5.9	275	2	H35216	FP31 protein - fow
116	106	6.3	1103	2	T22889	hypothetical prote	189	99	5.9	313	2	J01862	31R protein - vari
117	105.5	6.2	423	2	T29549	hypothetical prote	190	99	5.9	424	2	B36109	pregnancy-specific
118	105.5	6.2	582	1	BNRT3S	myelin-associated	191	99	5.9	458	1	RWHUT4	T-cell surface gly
119	105.5	6.2	626	1	BNRT3	myelin-associated	192	99	5.9	773	2	T46283	hypothetical prote
120	105.5	6.2	662	2	C40862	heparin-binding gr	193	99	5.9	812	1	A36477	fibroblast growth
121	105.5	6.2	729	2	A56795	fibroblast growth	194	99	5.9	818	2	JC4058	fibroblast growth
122	105.5	6.2	733	2	I49293	fibroblast growth	195	99	5.9	862	2	I49583	differentiation an
123	105.5	6.2	822	1	TVHUF6	fibroblast growth	196	99	5.9	1375	2	T13822	frazzled gene prot
124	105.5	6.2	822	2	I45289	fibroblast growth	197	99	5.9	1526	2	T13823	frazzled gene prot
125	105.5	6.2	822	2	S29840	fibroblast growth	198	98.5	5.8	288	2	T24824	hypothetical prote
126	105.5	6.2	940	2	A40985	projectin - fruit	199	98.5	5.8	602	2	A45769	acetylcholine rece
127	104.5	6.2	268	2	T23555	hypothetical prote	200	98.5	5.8	1021	2	T42634	connectin/titin -
128	104.5	6.2	392	2	B44194	poliovirus recepto	201	98.5	5.8	1330	2	S49100	embryonic receptor
129	104.5	6.2	417	2	A44194	poliovirus recepto	202	98.5	5.8	1333	2	I78875	receptor tyrosine
130	104.5	6.2	1184	2	T09484	cartilage intermed	203	98	5.8	880	1	JC4166	protein-tyrosine k
131	104.5	6.2	2222	2	T13924	sdk protein - frui	204	98	5.8	880	2	B53743	protein-tyrosine k
132	104	6.2	299	1	AHRB	ig alpha chain C r	205	98	5.8	1499	2	I50212	protein-tyrosine-p
133	104	6.2	338	2	S09276	ig alpha chain C r	206	97.5	5.8	144	2	S36308	T-cell receptor de
134	104	6.2	800	2	A48991	heparin-binding gr	207	97.5	5.8	354	1	S04243	proteoglycan link
135	104	6.2	801	2	I55363	fibroblast growth	208	97.5	5.8	476	1	A35104	brain-derived neur
136	104	6.2	882	2	I38912	receptor tyrosine	209	97.5	5.8	588	2	I37202	B-CAM protein - hu
137	103.5	6.1	628	2	I38000	Lutheran blood gro	210	97.5	5.8	729	2	A49120	fibroblast growth
138	103.5	6.1	819	1	TVCHFG	fibroblast growth	211	97.5	5.8	764	2	A49448	irregular chiasm C
139	103.5	6.1	822	1	TVM5FG	fibroblast growth	212	97.5	5.8	821	1	S06943	brain-derived neur
140	103.5	6.1	824	2	S36439	fibroblast growth	213	97.5	5.8	876	2	I49152	protein-tyrosine k
141	103.5	6.1	832	2	JH0393	fibroblast growth	214	97.5	5.8	1348	2	AH1115	cell surface prote
142	103.5	6.1	1356	2	JC1402	protein-tyrosine k	215	97.5	5.8	1443	2	I50600	neogenin - chicken
143	103.5	6.1	1612	2	T30805	duiti protein - mo	216	97	5.7	116	2	D24891	T-cell receptor al
144	103	6.1	811	2	A41054	fasciclin II, tran	217	97	5.7	307	2	S55596	hypothetical prote
145	103	6.1	873	2	B41054	fasciclin II pi-li	218	97	5.7	474	1	OMHUIB	alpha-1-B-glycopro
146	103	6.1	2783	2	T34416	hypothetical prote	219	97	5.7	739	2	A41288	vascular cell adhe
147	102.5	6.1	318	2	F72171	K9R protein - vari	220	97	5.7	751	2	T42597	DNA helicase/prima
148	102.5	6.1	1176	2	JN0583	myosin-light-chain	221	97	5.7	824	2	S24108	protein-tyrosine k
149	102.5	6.1	1651	2	T14160	transmembrane rece	222	97	5.7	1028	2	A53449	plasmacytoma-assoc
150	102	6.0	637	2	B33785	myelin-associated	223	97	5.7	1147	2	A59307	myosin-light-chain
151	102	6.0	702	2	A36319	carcinoembryonic a	224	97	5.7	1274	2	S55050	cardiac myosin-bin
152	102	6.0	800	1	TVHUF2F	fibroblast growth	225	97	5.7	1367	2	A41228	protein-tyrosine k
153	102	6.0	871	1	I48696	protein-tyrosine k	226	97	5.7	1535	2	S46224	peroxidase - frui
154	102	6.0	881	1	I48697	protein-tyrosine k	227	96.5	5.7	288	2	A31326	T-cell receptor de
155	101.5	6.0	392	1	RWHUPD	poliovirus recepto	228	96.5	5.7	313	2	T28598	hypothetical prote
156	101.5	6.0	413	2	S65948	hemolin - cecropia	229	96.5	5.7	772	2	T13078	KIAA0992 protein -
157	101.5	6.0	413	2	A37778	hemolin precursor	230	96.5	5.7	1200	2	T17404	hyalin - sea urchi
158	101.5	6.0	417	1	RWHUPA	poliovirus recepto	231	96.5	5.7	2013	2	AD1129	probable peptidogl
159	101.5	6.0	599	2	T16774	hypothetical prote	232	96	5.7	243	2	S25755	ig lambda chain -
160	101.5	6.0	898	2	A40114	fasciclin II precu	233	96	5.7	407	2	C82428	glucose-1-phosphat
161	101.5	6.0	1260	1	S05479	neural cell adhesi	234	96	5.7	462	2	I38404	neu differentiation
162	101	6.0	267	2	A38442	probable tumor sup	235	96	5.7	640	2	A43273	her regulin precurs
163	101	6.0	584	2	I50419	s-gicerin precurs	236	96	5.7	739	2	JN0581	vascular cell adhe
164	101	6.0	620	2	JH0593	Schwann cell myeli	237	96	5.7	1262	1	B48758	protein-tyrosine-p
165	101	6.0	797	2	S38579	fibroblast growth	238	96	5.7	1496	1	A48758	protein-tyrosine-p
166	101	6.0	868	2	A46512	CD22 homolog/B lym	239	96	5.7	1530	2	AH1396	peptidoglycan anch
167	101	6.0	890	1	A53743	protein-tyrosine k	240	96	5.7	6805	2	S20901	titin - rabbit (fr
168	101	6.0	1742	2	S24600	projectin - fruit	241	95.5	5.7	243	2	A53244	leukocyte antigen
169	101	6.0	6831	2	A88852	protein unc-22 [im	242	95.5	5.7	1132	2	A35089	myosin-binding pro
170	101	6.0	6839	2	S57242	twitchin [similar]	243	95	5.6	137	2	I46628	rearranged T-cell
171	101	6.0	7160	2	T27935	hypothetical prote	244	95	5.6	814	1	A39752	fibroblast growth
172	100.5	6.0	355	1	LKCH	proteoglycan link	245	95	5.6	994	2	I49276	c-mer tyrosine kin
173	100.5	6.0	408	1	LKRT2	proteoglycan link	246	95	5.6	1450	2	A44027	165k myofibrillar
174	100.5	6.0	647	2	A35648	B-cell adhesion pr	247	95	5.6	1501	2	I58148	protein-tyrosine-p
175	100.5	6.0	700	1	HYHUMB	meprin A (EC 3.4.2	248	95	5.6	1863	2	S46217	protein-tyrosine-p

249	95	5.6	1907	2	S50893	protein-tyrosine-p	322	89.5	5.3	142	2	S04664	T-cell receptor de
250	95	5.6	1945	2	T13937	plexin A - fruit f	323	89.5	5.3	157	2	S04915	T-cell receptor de
251	94.5	5.6	135	2	S36314	T-cell receptor	324	89.5	5.3	243	2	A37982	calcium vector pro
252	94.5	5.6	315	1	HNW24X	hemagglutinin prec	325	89.5	5.3	275	2	A28928	pregnancy-specific
253	94.5	5.6	328	2	S30444	ST2 protein - huma	326	89.5	5.3	531	2	S20900	tinin - mouse (fra
254	94.5	5.6	477	1	I73631	brain-derived neur	327	89.5	5.3	823	2	B35963	protein-tyrosine k
255	94.5	5.6	588	2	JH0506	adhesion molecule	328	89.5	5.3	829	1	IUHUCP	cadherin 3 precurs
256	94.5	5.6	588	2	A45254	surface glycoprote	329	89.5	5.3	1465	2	S43529	165K protein, skl
257	94.5	5.6	822	1	A56853	brain-derived neur	330	89.5	5.3	1483	2	C97012	probably celluloso
258	94.5	5.6	875	2	T33434	hypothetical prote	331	89	5.3	101	2	A39165	proteoglycan link
259	94.5	5.6	888	2	S23065	ufo protein - mous	332	89	5.3	145	2	I46629	rearranged T-cell
260	94.5	5.6	999	2	I38547	novel cellular pro	333	89	5.3	279	2	S04693	T-cell receptor de
261	94.5	5.6	1356	2	T03096	CD0 protein - rat	334	89	5.3	282	2	C28928	pregnancy-specific
262	94	5.6	118	2	I71934	MHC class II I-A-a	335	89	5.3	413	1	IKEC5A	colicin V secretio
263	94	5.6	268	2	A56446	Ig heavy chain V r	336	89	5.3	523	2	I50478	neurolin - goldfis
264	94	5.6	313	2	H36854	hemagglutinin - va	337	89	5.3	682	2	A35969	heparin-binding gr
265	94	5.6	422	2	S32357	glial growth facto	338	89	5.3	769	2	S16236	fibroblast growth
266	94	5.6	583	2	G84829	probable FRR2 fami	339	89	5.3	822	2	A45081	fibroblast growth
267	94	5.6	887	2	AD2009	hypothetical prote	340	89	5.3	822	2	A41794	keratinocyte growt
268	94	5.6	943	2	F69543	ATP-dependent RNA	341	89	5.3	1020	2	S05944	neural cell surf
269	94	5.6	1091	2	S01998	contactin precursor	342	88.5	5.2	524	2	S35341	kettin - fruit fly
270	94	5.6	1325	2	A64905	ydek protein - Esc	343	88.5	5.2	3562	2	A47171	chondroitin sulfat
271	93.5	5.5	142	2	S36307	T-cell receptor de	344	88	5.2	131	2	T20334	hypothetical prote
272	93.5	5.5	259	2	JC7109	ST2V protein - hum	345	88	5.2	247	2	A55717	myelin/oligodendro
273	93.5	5.5	329	1	A48754	B7-2 antigen - hum	346	88	5.2	568	2	A45804	Ig mu chain C regi
274	93.5	5.5	587	2	JH0464	DM-GRASP precursor	347	88	5.2	850	2	JC5700	Erbb kinase activa
275	93.5	5.5	799	2	S18209	fibroblast growth	348	88	5.2	1021	2	A57112	contactin precursor
276	93	5.5	432	1	RWC2T4	T-cell surface gly	349	88	5.2	1711	2	AB1283	peptidoglycan link
277	93	5.5	435	2	D33258	pregnancy-specific	350	88	5.2	2051	2	T30938	receptor tyrosine
278	93	5.5	1018	2	JC4211	neural adhesion pr	351	87.5	5.2	205	2	A48929	activated B-cell p
279	93	5.5	1643	2	T14274	versican precursor	352	87.5	5.2	246	2	A47712	myelin/oligodendro
280	93	5.5	3381	2	T42389	versican precursor	353	87.5	5.2	637	2	C43273	heregulin precursor
281	92.5	5.5	144	2	S36322	T-cell receptor de	354	87.5	5.2	742	2	J90326	alcohol dehydrogen
282	92.5	5.5	275	2	JC7604	CD86 spliced varia	355	87.5	5.2	757	2	I45956	polymeric immunogl
283	92.5	5.5	424	2	A34595	pregnancy-specific	356	87.5	5.2	1197	2	T30581	neural cell adhesi
284	92.5	5.5	713	2	S01028	fibroblast growth	357	87.5	5.2	1897	1	TDHULK	leukocyte antigen-
285	92.5	5.5	764	1	QRHUGS	secretory componen	358	87	5.2	139	2	S36302	T-cell receptor de
286	92.5	5.5	1321	2	T00382	hypothetical prote	359	87	5.2	364	2	A30521	myeloid cell surfa
287	92	5.5	116	2	E24891	T-cell receptor al	360	87	5.2	365	2	I72171	HLA-Aw33.1, HLA-Aw
288	92	5.5	132	2	A55410	Ig light chain V r	361	87	5.2	818	2	T19120	hypothetical prote
289	92	5.5	323	2	A48997	tumor surface anti	362	87	5.2	936	2	I40711	sagB protein - Cam
290	92	5.5	583	2	I39428	alcam - human	363	87	5.2	1451	2	S42167	190K protein - hum
291	92	5.5	639	2	I61719	neu differentiation	364	86.5	5.1	115	2	C24891	T-cell receptor al
292	92	5.5	829	2	JC4583	fibroblast growth	365	86.5	5.1	310	1	HNVZRA	hemagglutinin prec
293	92	5.5	1005	2	T18537	Ig heavy chain - c	366	86.5	5.1	363	2	JH0542	class I histocompa
294	92	5.5	13055	2	T16580	hypothetical prote	367	86.5	5.1	465	2	A82211	MurT/nudix family
295	91.5	5.4	240	2	JL0143	antigen BCM1 precu	368	86.5	5.1	894	1	A41527	protein-tyrosine k
296	91.5	5.4	330	2	A29915	teratocarcinoma gl	369	86.5	5.1	1348	2	S51656	vascular endotheli
297	91.5	5.4	576	2	A32604	interleukin-1 rece	370	86.5	5.1	1363	2	I58375	protein-tyrosine k
298	91.5	5.4	647	2	B41288	vascular cell adhe	371	86.5	5.1	1637	2	T46438	hypothetical prote
299	91.5	5.4	1241	2	T37190	nephlin - human	372	86.5	5.1	1894	2	C54689	protein-tyrosine-p
300	91.5	5.4	2013	2	A11489	probable peptidogl	373	86.5	5.1	2025	2	T21588	hypothetical prote
301	91	5.4	282	2	T17219	hypothetical prote	374	86.5	5.1	3157	2	B70969	probable PPE prote
302	90.5	5.4	309	2	S15674	cell surface glyco	375	86	5.1	116	2	B24891	T-cell receptor al
303	90.5	5.4	341	2	B95403	probable ABC trans	376	86	5.1	150	2	I46624	rearranged T-cell
304	90.5	5.4	361	2	PN0020	fibroblast growth	377	86	5.1	270	2	A34636	FC-gamma receptor
305	90.5	5.4	474	1	C39667	brain-derived neur	378	86	5.1	357	2	S09265	Ig alpha chain C r
306	90.5	5.4	476	1	B39667	brain-derived neur	379	86	5.1	398	2	S17428	interleukin-1 rece
307	90.5	5.4	487	2	S65133	butyrophilin - mou	380	86	5.1	421	2	T46266	hypothetical prote
308	90.5	5.4	631	1	A48346	cell fusion glycop	381	86	5.1	480	2	B56182	fibroblast growth
309	90.5	5.4	631	1	VGNZPD	cell fusion glycop	382	86	5.1	497	2	JC2456	pyruvate kinase (B
310	90.5	5.4	821	1	A39667	brain-derived neur	383	86	5.1	964	2	T15746	hypothetical prote
311	90.5	5.4	860	2	JC5702	Erbb kinase activa	384	86	5.1	975	1	TVMSKT	protein-tyrosine k
312	90.5	5.4	868	2	JC5701	Erbb kinase activa	385	86	5.1	1327	2	T09402	immunoglobulin-lik
313	90	5.3	320	2	S12839	Ig heavy chain pre	386	86	5.1	1345	2	H90975	hypothetical prote
314	90	5.3	526	2	T27581	hypothetical prote	387	86	5.1	1417	2	H90670	probable invasin l
315	90	5.3	789	2	T28714	hypothetical prote	388	86	5.1	1417	2	D85521	probable adhesin e
316	90	5.3	821	1	TVHUF2	fibroblast growth	389	86	5.1	2660	2	E85822	probable invasin Z
317	90	5.3	1018	2	A54744	contactin 1 precur	390	85.5	5.1	210	2	A56169	Ig kappa chain V r
318	90	5.3	1240	2	T03097	CD0 protein - huma	391	85.5	5.1	216	2	J50245	Ig lambda chain NI
319	90	5.3	1343	2	E90893	hypothetical prote	392	85.5	5.1	218	2	S68241	Ig kappa chain V r
320	90	5.3	1343	2	D85724	hypothetical prote	393	85.5	5.1	302	2	C36464	fibroblast growth
321	90	5.3	1355	2	T28715	hypothetical prote	394	85.5	5.1	329	2	A44065	fibroblast growth

395	85.5	5.1	330	2	I46691	CD86 precursor - r	468	83.5	4.9	372	2	C39371	lg V-region-like B
396	85.5	5.1	332	2	C83610	sulfate-binding pr	469	83.5	4.9	496	2	S60685	envelope protein E
397	85.5	5.1	388	2	B48899	beta-lactamase [BC	470	83.5	4.9	498	2	S11246	IAG-3 protein prec
398	85.5	5.1	397	2	C43354	pregnancy-specific	471	83.5	4.9	527	2	D75127	hypothetical prote
399	85.5	5.1	419	2	JC4123	pregnancy-specific	472	83.5	4.9	629	2	AE1525	probable peptidogl
400	85.5	5.1	627	2	S14683	ig mu chain precu	473	83.5	4.9	705	2	S51635	fibroblast growth
401	85.5	5.1	636	2	I61718	neu differentiation	474	83.5	4.9	707	2	A38429	keratinocyte growt
402	85.5	5.1	869	2	A85983	conserved hypotet	475	83.5	4.9	748	2	S41050	fibroblast growth
403	85.5	5.1	869	2	S72760	ppsl protein - Myc	476	83.5	4.9	750	2	S41051	fibroblast growth
404	85.5	5.1	1094	2	S22573	DNA-directed DNA p	477	83.5	4.9	911	2	I56552	synapse-associated
405	85.5	5.1	1518	2	S37928	probable purine nu	478	83.5	4.9	987	2	A88746	protein C18F3.2 [i
406	85.5	5.1	1691	1	D54689	protein-tyrosine-p	479	83.5	4.9	1002	2	T19226	hypothetical prote
407	85.5	5.1	2541	2	T29340	hypothetical prote	480	83.5	4.9	1028	2	C88364	protein C13B4.1 li
408	85	5.0	104	2	S07705	T-cell receptor al	481	83.5	4.9	1287	2	T30988	hypothetical prote
409	85	5.0	137	2	S36303	T-cell receptor de	482	83.5	4.9	3898	1	GNWVHC	genome polyprotein
410	85	5.0	166	2	A33402	pregnancy-specific	483	83	4.9	115	2	B32071	T-cell receptor de
411	85	5.0	247	2	S58394	myelin/oligodendro	484	83	4.9	233	2	S25747	ig lambda chain -
412	85	5.0	265	2	A55811	carcinoembryonic a	485	83	4.9	422	2	B95001	conserved domain p
413	85	5.0	273	2	B28928	pregnancy-specific	486	83	4.9	428	2	A37658	pregnancy-specific
414	85	5.0	428	2	I57486	pregnancy-specific	487	83	4.9	445	2	A97873	conserved hypotet
415	85	5.0	428	2	JS0032	pregnancy-specific	488	83	4.9	977	2	S49004	tyrosine kinase Mp
416	85	5.0	480	2	A56182	fibroblast growth	489	83	4.9	1151	2	A45226	integrin alpha-1 c
417	85	5.0	496	2	PQ0666	envelope protein -	490	83	4.9	3029	2	S76109	hypothetical prote
418	85	5.0	769	1	QRRTGS	secretory componen	491	82.5	4.9	185	2	S37479	T-cell receptor al
419	85	5.0	776	2	S41628	genome polyprotein	492	82.5	4.9	218	2	JC5810	monoclonal antibod
420	85	5.0	904	2	I38757	homolog of Drosoph	493	82.5	4.9	218	2	B47712	myelin/oligodendro
421	85	5.0	913	1	A47543	R-cadherin precurs	494	82.5	4.9	229	2	A46527	B-cell-specific me
422	85	5.0	926	2	I38756	homolog of Drosoph	495	82.5	4.9	230	2	A56210	neu differentiation
423	85	5.0	1005	2	A42265	alpha-mannosidase	496	82.5	4.9	316	2	C37028	MHC class I histoc
424	85	5.0	2325	2	A61208	chondroitin sulfat	497	82.5	4.9	353	2	S51242	heparin-binding fi
425	85	5.0	2409	1	A60979	versican precursor	498	82.5	4.9	355	2	I80169	class I histocoma
426	84.5	5.0	241	2	S32359	glial growth facto	499	82.5	4.9	356	1	S55437	translational releas
427	84.5	5.0	266	2	T29613	hypothetical prote	500	82.5	4.9	395	2	D43354	pregnancy-specific
428	84.5	5.0	286	2	A28333	carcinoembryonic a	501	82.5	4.9	417	2	B83277	pregnancy-specific
429	84.5	5.0	419	2	A33258	pregnancy-specific	502	82.5	4.9	453	2	B83380	hypothetical prote
430	84.5	5.0	419	2	A31135	pregnancy-specific	503	82.5	4.9	645	2	B43273	heregulin, splice
431	84.5	5.0	432	2	S30193	T-cell surface gly	504	82.5	4.9	651	2	A96781	unknown protein P9
432	84.5	5.0	436	2	E69371	bile acid-inducibl	505	82.5	4.9	662	2	I61722	neu differentiatio
433	84.5	5.0	457	2	A27449	T-cell surface gly	506	82.5	4.9	757	1	S48841	secretory componen
434	84.5	5.0	474	2	S15590	ig heavy chain - h	507	82.5	4.9	793	2	S59067	penton long fiber
435	84.5	5.0	568	2	I58106	gene DMR-N9 protei	508	82.5	4.9	822	2	B19947	fibroblast growth
436	84.5	5.0	642	2	E72215	oligopeptide ABC t	509	82.5	4.9	822	2	B49151	fibroblast growth
437	84.5	5.0	795	2	T12487	hypothetical prote	510	82.5	4.9	942	2	S23251	protein-tyrosine k
438	84.5	5.0	821	1	TVM5BK	fibroblast growth	511	82.5	4.9	975	2	T42576	probable envelope
439	84.5	5.0	941	1	TVMVMD	protein-tyrosine k	512	82.5	4.9	1015	2	T32186	hypothetical prote
440	84.5	5.0	980	1	TVC7MD	macrophage colony-	513	82.5	4.9	1648	2	S61654	probable membrane
441	84.5	5.0	1142	2	S36845	myosin-binding pro	514	82	4.9	147	2	I46625	rearranged T-cell
442	84.5	5.0	1439	2	A11476	cell surface prote	515	82	4.9	150	2	S36312	T-cell receptor de
443	84.5	5.0	1433	2	T30261	chitinase (EC 3.2.	516	82	4.9	242	2	S06942	hypothetical prote
444	84.5	5.0	2508	2	S61441	surface-associated	517	82	4.9	250	2	H81854	probable zinc-bind
445	84	5.0	238	2	C90909	probable major tai	518	82	4.9	261	2	C85681	hypothetical prote
446	84	5.0	255	2	JC7593	SH2 domain-contain	519	82	4.9	324	2	G43354	pregnancy-specific
447	84	5.0	315	1	HNW2VT	hemagglutinin prec	520	82	4.9	326	2	F43354	pregnancy-specific
448	84	5.0	315	2	T37438	hemagglutinin - va	521	82	4.9	333	2	A43354	pregnancy-specific
449	84	5.0	321	2	D39371	ig V-region-like B	522	82	4.9	335	2	A33514	pregnancy-specific
450	84	5.0	406	2	E43354	pregnancy-specific	523	82	4.9	335	2	H43354	ig V-region-like B
451	84	5.0	452	1	MHRU	ig mu chain C regi	524	82	4.9	340	2	T28137	ig alpha chain C r
452	84	5.0	453	2	S37768	ig mu chain C regi	525	82	4.9	352	2	S09266	pregnancy-specific
453	84	5.0	473	1	MHHUM	ig mu chain C regi	526	82	4.9	426	2	C55181	pregnancy-specific
454	84	5.0	621	2	B57431	myosin-binding C-p	527	82	4.9	426	2	S09016	pregnancy-specific
455	84	5.0	781	2	S51592	XyNB precursor - R	528	82	4.9	426	2	B35334	pregnancy-specific
456	84	5.0	939	2	AF2503	hypothetical prote	529	82	4.9	436	2	B55181	pregnancy-specific
457	84	5.0	1065	2	H95321	NolG efflux transp	530	82	4.9	473	2	T35997	conserved hypotet
458	84	5.0	1912	2	A56178	protein-tyrosine-p	531	82	4.9	1086	2	AF1662	cellobiose-phospho
459	84	5.0	2397	1	A55535	versican precursor	532	82	4.9	1160	2	F88369	protein unc-52 [im
460	83.5	4.9	264	2	F27579	T-cell receptor al	533	82	4.9	1385	2	D89824	hypothetical prote
461	83.5	4.9	275	1	RWHUAC	T-cell receptor al	534	82	4.9	1898	2	S46216	leukocyte antigen-
462	83.5	4.9	288	2	A45803	B-cell-restricted	535	81.5	4.8	85	2	S08109	carcinoembryonic a
463	83.5	4.9	317	2	JO1118	Fc gamma (IgG) rec	536	81.5	4.8	140	2	C24747	T-cell receptor be
464	83.5	4.9	329	2	AG0010	exported sulfate-b	537	81.5	4.8	213	2	S68213	ig heavy chain (Na
465	83.5	4.9	339	2	S09264	ig alpha chain C r	538	81.5	4.8	235	2	I50610	T-cell surface gly
466	83.5	4.9	344	1	RWRTC2	T-cell surface gly	539	81.5	4.8	314	1	HNWZVM	hemagglutinin prec
467	83.5	4.9	363	2	S07113	class I histocoma	540	81.5	4.8	314	1	JQ1793	hemagglutinin prec

541	81.5	4.8	339	2	T28138	Ig V-region-like B	614	79.5	4.7	341	2	S16521	mitosis-specific c
542	81.5	4.8	426	1	JC15964	pregnancy-specific	615	79.5	4.7	360	2	S09271	Ig alpha chain C r
543	81.5	4.8	650	1	JC1450	fibroblast growth	616	79.5	4.7	361	2	S68089	actin 2 - Arabidop
544	81.5	4.8	687	2	A49636	soluble vascular e	617	79.5	4.7	474	2	B82227	exodeoxyribonuclea
545	81.5	4.8	699	2	A12686	proteinase II [imp	618	79.5	4.7	483	2	A97744	isocitrate dehydro
546	81.5	4.8	699	2	E97468	dipeptidyl aminope	619	79.5	4.7	495	2	A55181	pregnancy-specific
547	81.5	4.8	707	2	A54846	fibroblast growth	620	79.5	4.7	610	2	A55939	diacylglycerol S
548	81.5	4.8	822	2	B54846	fibroblast growth	621	79.5	4.7	637	2	T49099	hypothetical prote
549	81.5	4.8	1015	2	T32984	hypothetical prote	622	79.5	4.7	701	2	T17243	endo-beta-N-acetyl
550	81.5	4.8	1462	1	A69809	probable multifunc	623	79.5	4.7	878	2	A83748	starch phosphoryla
551	81.5	4.8	3507	2	T34513	hypothetical prote	624	79.5	4.7	974	2	S34189	homeodomain-intera
552	81	4.8	117	2	A49883	T-cell receptor alp	625	79.5	4.7	1209	2	T14357	neural cell adhesi
553	81	4.8	139	2	I46630	rearranged T-cell	626	79.5	4.7	1265	1	A37967	complement C3 prec
554	81	4.8	145	2	I46631	rearranged T-cell	627	79.5	4.7	1663	1	C3RT	Ig lambda chain pr
555	81	4.8	206	2	A46305	biliary glycoprote	628	79	4.7	118	2	S04519	pregnancy-specific
556	81	4.8	315	1	HNZVTV	hemagglutinin prec	629	79	4.7	132	2	P00060	heretulin precursor
557	81	4.8	381	1	I51174	Ig heavy chain - R	630	79	4.7	210	2	JC4122	T-cell receptor be
558	81	4.8	402	2	T09062	probable advanced	631	79	4.7	241	2	D43273	hypothetical prote
559	81	4.8	432	1	RWQGT4	T-cell surface gly	632	79	4.7	345	2	A46052	vascular cell adhe
560	81	4.8	438	2	E96545	hypothetical prote	633	79	4.7	370	2	S29139	aggregran - pig (fr
561	81	4.8	449	2	A45557	matrix, capsid, nu	634	79	4.7	374	2	A46352	ORF1 protein - Chl
562	81	4.8	450	2	S25162	gag protein - fel	635	79	4.7	407	2	T08732	hypothetical prote
563	81	4.8	509	2	JC6203	SP8 binding protei	636	79	4.7	451	2	S71754	cellular hepatitis
564	81	4.8	767	2	F86383	hypothetical prote	637	79	4.7	502	2	T02746	cyclin A-like prot
565	81	4.8	823	2	S44873	ZC21.2 protein - C	638	79	4.7	567	2	S29498	lymphocyte antigen
566	81	4.8	1433	1	A36734	protein J - Yersin	639	79	4.7	716	1	WZBER6	77.8K DNA helicase
567	81	4.8	1492	2	T14652	phage lambda-relat	640	79	4.7	741	2	E83265	probable two-compo
568	81	4.8	1545	2	T14966	phage lambda-relat	641	79	4.7	742	2	P84643	hypothetical prote
569	81	4.8	1650	2	S53457	dominant autoantig	642	79	4.7	1255	2	T06267	nematodes resistan
570	80.5	4.8	133	2	S57870	T cell receptor Ck	643	79	4.7	1298	2	A48999	protein-tyrosine k
571	80.5	4.8	142	2	B28344	VpreB protein prec	644	79	4.7	1315	2	T28679	chromodomain helic
572	80.5	4.8	366	2	T37523	MHC class I histoc	645	79	4.7	1388	2	T38720	genome polypeptid
573	80.5	4.8	426	2	B33258	pregnancy-specific	646	79	4.7	3414	1	GNWVNE	T-cell receptor be
574	80.5	4.8	426	2	A53341	pregnancy-specific	647	79	4.7	136	2	J00473	T-cell receptor de
575	80.5	4.8	441	2	T31482	hypothetical prote	648	78.5	4.7	136	2	S36320	integrin-associate
576	80.5	4.8	551	2	A30342	interleukin-2 rece	649	78.5	4.7	215	2	A57843	inositol monophosp
577	80.5	4.8	820	2	S17295	fibroblast growth	650	78.5	4.7	230	2	S33161	class I histocompa
578	80.5	4.8	874	2	T29548	hypothetical prote	651	78.5	4.7	269	2	S57494	MHC class I histoc
579	80.5	4.8	1216	2	S60613	protein-tyrosine-p	652	78.5	4.7	270	2	S65739	MHC class I histoc
580	80.5	4.8	1493	2	T10757	MAP kinase kinase	653	78.5	4.7	279	2	S53363	MHC class I histoc
581	80.5	4.8	1641	2	D82704	conserved hypotet	654	78.5	4.7	324	2	S36646	MHC class I histoc
582	80.5	4.8	2482	2	I48922	cation-independent	655	78.5	4.7	335	2	A75363	conserved hypotet
583	80.5	4.8	2483	1	A49617	insulin-like growt	656	78.5	4.7	355	2	I80171	pregnancy-specific
584	80	4.7	124	2	F83560	probable type II s	657	78.5	4.7	363	2	S03537	Ig heavy chain pre
585	80	4.7	133	2	I46632	rearranged T-cell	658	78.5	4.7	366	2	I37544	E-selectin precurs
586	80	4.7	147	2	S25499	T-cell receptor al	659	78.5	4.7	366	2	S42823	DNA ligase (ATP) (
587	80	4.7	194	2	T29925	hypothetical prote	660	78.5	4.7	366	2	B37028	probable lipoprote
588	80	4.7	218	2	A36040	Ig heavy chain V-I	661	78.5	4.7	407	2	C83589	cadherin 4 precurs
589	80	4.7	287	2	S71192	mitosis-specific c	662	78.5	4.7	419	2	B54312	probable tenascin
590	80	4.7	304	1	RWCHH7	cell surface glyco	663	78.5	4.7	568	2	A34891	hypothetical prote
591	80	4.7	365	2	A45847	MHC class I histoc	664	78.5	4.7	612	2	B42755	sex factor aggrega
592	80	4.7	549	2	S04845	Ig heavy chain pre	665	78.5	4.7	624	2	A29066	probable membrane
593	80	4.7	684	2	F97148	aldehyde:ferredoxi	666	78.5	4.7	768	2	A29066	hyperplastic discs
594	80	4.7	707	2	JC7763	neuronal leucine-r	667	78.5	4.7	787	2	S73873	probable tenascin
595	80	4.7	748	2	T37097	probable secreted	668	78.5	4.7	916	2	C38992	hypothetical prote
596	80	4.7	893	2	A37084	surface-array prot	669	78.5	4.7	1059	2	T22545	hypothetical prote
597	80	4.7	939	2	AE2275	hypothetical prote	670	78.5	4.7	1188	1	A57064	protein-tyrosine-p
598	80	4.7	999	2	S72267	Ca2+-transporting	671	78.5	4.7	1243	2	S60138	sex factor aggrega
599	80	4.7	1123	2	S36846	myosin-binding pro	672	78.5	4.7	1609	2	S25345	probable membrane
600	80	4.7	1582	2	T15308	hypothetical prote	673	78.5	4.7	2895	2	T08437	hyperplastic discs
601	80	4.7	2481	2	A43908	fibronectin - Afri	674	78	4.6	4006	2	T09070	probable tenascin
602	80	4.7	3488	2	T34418	T-cell receptor de	675	78	4.6	121	2	A29080	hypothetical prote
603	79.5	4.7	104	2	S36305	T-cell receptor de	676	78	4.6	309	2	I49503	B-lymphocyte activ
604	79.5	4.7	114	2	JH0341	T-cell receptor al	677	78	4.6	335	2	C54312	pregnancy-specific
605	79.5	4.7	135	2	S38298	T-cell receptor de	678	78	4.6	336	2	S42632	Fit-1S protein pre
606	79.5	4.7	137	2	C45893	T-cell receptor al	679	78	4.6	342	2	E71101	probable geranylge
607	79.5	4.7	140	2	I46638	rearranged T-cell	680	78	4.6	367	2	A83303	putrescine-binding
608	79.5	4.7	147	2	I46623	Ig lambda chain -	681	78	4.6	508	2	A33378	fasciclin III prec
609	79.5	4.7	232	2	S25756	fibroblast growth	682	78	4.6	592	2	S25705	Ig mu chain - shee
610	79.5	4.7	254	2	G00031	B7 protein - red-c	683	78	4.6	617	2	E64734	novel antigen rece
611	79.5	4.7	289	2	G00031	MHC ChIA chain - c	684	78	4.6	684	2	S60266	killer cell inhibi
612	79.5	4.7	313	2	I36958	Ig alpha chain C r	685	78	4.6	841	2	JC5894	glycoprotein 350/2
613	79.5	4.7	335	2	S09275		686	78	4.6	886	2	S29605	

687	78	4.6	1185	2	A42404	collagen adhesin -	760	76.5	4.5	771	2	A83348	probable aldehyde
688	78	4.6	1203	2	T04294	hypothetical prote	761	76.5	4.5	874	2	C84513	Mutator-like trans
689	78	4.6	1273	1	T0RPLT	leukocyte common a	762	76.5	4.5	1110	2	T19673	hypothetical prote
690	78	4.6	3848	2	T17414	TiPC protein - ali	763	76.5	4.5	1138	2	S24614	myosin-binding pro
691	78	4.6	4116	2	T13719	calo protein - fru	764	76.5	4.5	1350	2	AF2005	RNA polymerase bet
692	77.5	4.6	103	2	S36067	ig lambda chain -	765	76.5	4.5	1365	2	T30198	alkaline phosphata
693	77.5	4.6	104	2	G24402	T-cell receptor al	766	76.5	4.5	1832	2	T31113	mucin-like glycopr
694	77.5	4.6	122	2	S16439	ig lambda chain -	767	76	4.5	120	2	C26945	T-cell receptor de
695	77.5	4.6	134	2	A45893	T-cell receptor al	768	76	4.5	133	2	P24402	T-cell receptor al
696	77.5	4.6	182	2	I83053	pregnancy-specific	769	76	4.5	141	2	S36318	T-cell receptor de
697	77.5	4.6	226	2	S25745	ig lambda chain -	770	76	4.5	145	2	S25743	ig lambda chain -
698	77.5	4.6	227	2	A33937	ig light chain (13	771	76	4.5	149	2	S36317	T-cell receptor de
699	77.5	4.6	235	2	S25758	ig lambda chain -	772	76	4.5	323	2	S06946	Fc gamma (IgG) rec
700	77.5	4.6	240	2	JC4121	pregnancy-specific	773	76	4.5	341	2	S26686	cAMP response elem
701	77.5	4.6	288	2	S29690	ig heavy chain VDJ	774	76	4.5	360	2	A27638	MHC class I histoc
702	77.5	4.6	295	2	A37412	T-cell receptor de	775	76	4.5	362	2	I72755	HLA-B*5602 - human
703	77.5	4.6	332	2	JN0067	pregnancy-specific	776	76	4.5	362	2	I72754	HLA-B*5601 - human
704	77.5	4.6	363	1	S42102	MHC class I histoc	777	76	4.5	365	2	I38439	MHC class I histoc
705	77.5	4.6	364	2	S59931	glycerophosphodie	778	76	4.5	366	2	I81231	lymphocyte antigen
706	77.5	4.6	364	2	S59932	glycerophosphodie	779	76	4.5	400	2	AD3364	probable acyl-CoA
707	77.5	4.6	364	2	S59933	glycerophosphodie	780	76	4.5	401	2	AB3552	branched-chain ami
708	77.5	4.6	364	2	A43576	glycerophosphodie	781	76	4.5	415	2	AB3637	alpha-methylacyl-C
709	77.5	4.6	364	2	S59934	glycerophosphodie	782	76	4.5	441	2	A96817	hypothetical prote
710	77.5	4.6	420	2	H84182	hypothetical prote	783	76	4.5	443	2	A96818	hypothetical prote
711	77.5	4.6	496	2	S60686	envelope protein E	784	76	4.5	446	2	F84451	probable uridylyl
712	77.5	4.6	509	2	S17597	ig delta chain (WI	785	76	4.5	457	1	RWMS74	T-cell surface gly
713	77.5	4.6	537	2	A46611	myosin-binding pro	786	76	4.5	478	2	AF1758	chitinase and chit
714	77.5	4.6	542	2	S45219	fiber protein - ca	787	76	4.5	496	2	FC4408	protein E - Tick-b
715	77.5	4.6	662	2	T16525	hypothetical prote	788	76	4.5	562	2	A65166	hypothetical 63.2K
716	77.5	4.6	704	2	A48040	meprin A (EC 3.4.2	789	76	4.5	562	2	B91194	probable enzyme [i
717	77.5	4.6	1009	2	T18533	CryIAC toxin-bind	790	76	4.5	562	2	C86041	probable enzyme yi
718	77.5	4.6	1036	2	B69368	hypothetical prote	791	76	4.5	609	2	T16135	hypothetical prote
719	77.5	4.6	1047	2	T41343	probable translati	792	76	4.5	649	2	T32755	hypothetical prote
720	77.5	4.6	1141	2	E89824	hypothetical prote	793	76	4.5	800	2	S37387	internalin A precu
721	77.5	4.6	1997	1	S12050	protein-tyrosine-p	794	76	4.5	959	1	P2XR13	outer capsid prote
722	77	4.6	95	2	D95201	hypothetical prote	795	76	4.5	1032	2	T34433	hypothetical prote
723	77	4.6	113	2	JH0340	T-cell receptor al	796	76	4.5	146	2	H32536	procollagen type V
724	77	4.6	135	2	S26512	T-cell receptor al	797	76	4.5	2232	2	T34434	hypothetical prote
725	77	4.6	146	2	S36323	T-cell receptor de	798	76	4.5	3716	2	E70969	probable PPE prote
726	77	4.6	309	2	T15747	hypothetical prote	799	76	4.5	4660	2	T42737	gp330 protein prec
727	77	4.6	331	2	T31517	hypothetical prote	800	75.5	4.5	116	2	S22558	ig heavy chain v r
728	77	4.6	356	2	F71624	refin PF80060w - m	801	75.5	4.5	146	2	H32536	T-cell receptor al
729	77	4.6	396	2	T45033	hypothetical prote	802	75.5	4.5	275	2	PS0402	basigin type III -
730	77	4.6	400	1	A39822	leukosialin precu	803	75.5	4.5	283	2	F87183	probable exported
731	77	4.6	403	2	I52590	m33-B isoform - mo	804	75.5	4.5	297	2	AG2955	hypothetical prote
732	77	4.6	430	2	T28143	tapasin 1 homolog,	805	75.5	4.5	297	2	F98327	hypothetical prote
733	77	4.6	433	2	T17289	hypothetical prote	806	75.5	4.5	312	2	H64560	probable tetraacyl
734	77	4.6	462	2	A41158	dipeptidyl-peptida	807	75.5	4.5	342	2	A45966	ig alpha chain C r
735	77	4.6	496	2	A43383	envelope glycoprot	808	75.5	4.5	366	2	T37527	MHC class I histoc
736	77	4.6	590	2	I56526	interleukin 1 rece	809	75.5	4.5	371	2	A53908	brevican precursor
737	77	4.6	630	2	T51281	beta-D-glucan exoh	810	75.5	4.5	398	2	A39371	ig V-region-like B
738	77	4.6	639	2	T20772	hypothetical prote	811	75.5	4.5	484	2	B64481	hypothetical prote
739	77	4.6	661	2	B54078	methyl-accepting c	812	75.5	4.5	549	2	G9618	complement subcomp
740	77	4.6	680	2	JC5895	killer cell inhibi	813	75.5	4.5	607	2	A95122	Tn5252, relaxase [
741	77	4.6	690	2	G84638	hypothetical prote	814	75.5	4.5	705	1	C1HURB	alcohol dehydrogen
742	77	4.6	800	2	AB1129	internalin A [impo	815	75.5	4.5	742	2	A49340	AAA family ATPase
743	77	4.6	846	2	T19179	hypothetical prote	816	75.5	4.5	769	2	E90158	hypothetical prote
744	77	4.6	1666	2	T38393	clathrin heavy cha	817	75.5	4.5	796	2	T20393	neurotrophic recep
745	77	4.6	2491	1	A28372	insulin-like growt	818	75.5	4.5	937	2	A45082	exodeoxyribonuclea
746	76.5	4.5	114	2	I38316	T-cell receptor be	819	75.5	4.5	1042	2	AF0739	hypothetical prote
747	76.5	4.5	115	2	S03513	T-cell receptor be	820	75.5	4.5	1294	2	T04278	hypothetical prote
748	76.5	4.5	120	2	S36306	T-cell receptor de	821	75.5	4.5	1662	2	T01893	hypothetical prote
749	76.5	4.5	151	2	I46626	rearranged T-cell	822	75.5	4.5	1902	2	C97702	cell surface antiq
750	76.5	4.5	239	2	G97165	flagellar hook ass	823	75	4.4	132	1	RWMSAV	T-cell receptor al
751	76.5	4.5	240	2	S01299	OX-45 membrane gly	824	75	4.4	141	2	S38389	T-cell receptor be
752	76.5	4.5	240	2	A39016	T-cell surface gly	825	75	4.4	233	2	S25744	ig kappa chain pre
753	76.5	4.5	255	2	JQ1240	hypothetical 29.3K	826	75	4.4	234	2	S01320	major tail protein
754	76.5	4.5	296	2	B39581	GRESAG protein 9u	827	75	4.4	238	2	D90876	probable Rieske pr
755	76.5	4.5	318	2	H75157	daunorubicin resis	828	75	4.4	252	2	H72554	probable tail comp
756	76.5	4.5	336	2	I48471	Fc gamma (IgG) rec	829	75	4.4	262	2	C85692	probable proteinas
757	76.5	4.5	477	2	H75026	oligopeptide abc t	830	75	4.4	301	2	AC3476	probable cross-
758	76.5	4.5	530	2	T18596	hypothetical prote	831	75	4.4	335	2	B33251	non-specific cross-
759	76.5	4.5	742	2	S38093	probable purine nu	832	75	4.4	346	2	C81088	alcohol dehydrogen

QY 283 PVF 285
DB 394 PVF 396

RESULT 5

S18252
heparan sulfate proteoglycan - mouse
N:Alternate names: perlecan
C:Species: Mus musculus (house mouse)
C>Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 12-Jul-2004
C:Accession: S18252; A31917; B31917; S66460
R:Noonan, D.M.; Fullie, A.; Valence, P.; Cai, S.; Horigan, E.; Sasaki, M.; Yamada, Y.; Hasegawa, J. Biol. Chem. 266, 22939-22947, 1991
A:Title: The complete sequence of perlecan, a basement membrane heparan sulfate proteoglycan molecule.
A:Reference number: S18252; MUID:92078153; PMID:1744087
A:Accession: S18252
A:Molecule type: mRNA
A:Residues: 1-3707 <NOO>
A:Cross-references: UNIPROT:Q05793; EMBL:M77174; NID:G200295; PIDN:AAA39911.1; PID:G200295
R:Noonan, D.M.; Horigan, E.A.; Ledbetter, S.R.; Vogeli, G.; Sasaki, M.; Yamada, Y.; Hasegawa, J. Biol. Chem. 263, 16379-16387, 1988
A:Title: Identification of cDNA clones encoding different domains of the basement membrane proteoglycan molecule.
A:Reference number: A92680; MUID:89034110; PMID:2972708
A:Accession: A31917
A:Molecule type: mRNA
A:Residues: 940-1601 <NO2>
A:Cross-references: GB:J04054; NID:G200252; PIDN:AAA39899.1; PID:G200253
A:Accession: B31917
A:Molecule type: mRNA
A:Residues: 1870-2600 <NO3>
A:Cross-references: GB:J04055; NID:G200300; PIDN:AAA39912.1; PID:G200301
R:Schulze, B.; Mann, K.; Battistutta, R.; Wiedemann, H.; Timpl, R. Eur. J. Biochem. 231, 551-556, 1995
A:Title: Structural properties of recombinant domain III-3 of perlecan containing a globular domain.
A:Reference number: S66460; MUID:95377282; PMID:7649154
A:Accession: S66460
A:Molecule type: protein
A:Residues: 1272-1274, 'X', 1276, 'X', 1278-1279 <SCH>
C:Keywords: glycoprotein
F:199-234/Domain: LDL receptor ligand-binding repeat homology <LDL1>
F:288-319/Domain: LDL receptor ligand-binding repeat homology <LDL2>
F:325-359/Domain: LDL receptor ligand-binding repeat homology <LDL3>
F:368-403/Domain: LDL receptor ligand-binding repeat homology <LDL4>
F:764-811/Domain: laminin-type EGF-like homology <LEG>
F:1159-1206/Domain: laminin-type EGF-like homology <LEG7>
F:1563-1610/Domain: laminin-type EGF-like homology <LEG8>
F:1613-1668/Domain: laminin-type EGF-like homology <LEG9>
F:3163-3198/Domain: EGF homology <EGF>
F:3270-3423/Domain: laminin G repeat homology <LG2>
F:3464-3492/Domain: EGF homology <EGF7>
F:1256, 1891, 2336, 2394, 2427/Binding site: carbohydrate (Asn) (covalent) #status predicted

Query Match 8.3%; Score 139.5; DB 2; Length 3707;
Best Local Similarity 25.1%; Pred. No. 0.055;
Matches 66; Conservative 31; Mismatches 105; Indels 61; Gaps 14;
QY 13 LTVDT-----YGRPILVEPESVTGPKWGDVNLPCYDPLQGYTQVLVWLVQSGDPVTIF 68
DB 2521 LTVDTGTVAPGPPQVVEES-----ELTLEA-----GHTATL--HCSATGNPPPT-- 2563
QY 69 LRDSGDHIQAKYQGRHLVSHKVPQGVDSLSQLSTLEMDRSHYTCVWTQPDGQVVRD 128
DB 2564 -----IHWKLRAPLPWQRIEGLN-TLVPRAVQDQSGQYICNAT-----NSAGHT 2608
QY 129 KTELRLVQKLSVKPTVTGSGYGTVPQ-----GMRISLQCGARGSPPIYSIYWKQOT 182
DB 2609 EATV-----LHVESPP-----YATIIPEHTSAQGNLVQLCLAHGTPPLTYQW--SLV 2656
QY 183 NNQEPKIVATLSTIL-FKPAVIADSGSYFCTAKGVSEQHSIDIVKVFVVKSSKLLTKTKT 241
DB 2657 GGVLPKAVVRNQLRLLEPTPEDSGRYCQVSNRVGSAE--AFAQVLVQSSSNLPDTS 2714

QY 242 ----EAPTTMTYPLKATSTVKQS 260
DB 2715 IPGSGTPTVQVTPQLETRNIGAS 2737

RESULT 6

PN0568
connectin 3B - chicken (fragment)
N:Alternate names: Cn3B protein
C:Species: Gallus gallus (chicken)
C>Date: 03-May-1994 #sequence_revision 07-Oct-1994 #text_change 09-Jul-2004
C:Accession: PN0568
R:Maruyama, K.; Endo, T.; Kume, H.; Kawamura, Y.; Kanzawa, N.; Nakauchi, Y.; Kimura, S.; Biochem. Biophys. Res. Commun. 194, 1288-1291, 1993
A:Title: A novel domain sequence of connectin localized at the 1 band of skeletal muscle
A:Reference number: PN0568; MUID:93356802; PMID:8352787
A:Accession: PN0568
A:Molecule type: mRNA
A:Residues: 1-1323 <MAR>
A:Cross-references: UNIPROT:Q08476; DBJ:D16541; NID:G391629; PID:D1004495; PID:G391630
A:Experimental source: skeletal muscle
C:Comment: This protein string-like single molecule spans from the Z line to the M line
Query Match 8.2%; Score 138.5; DB 2; Length 1323;
Best Local Similarity 21.4%; Pred. No. 0.018;
Matches 61; Conservative 38; Mismatches 117; Indels 69; Gaps 10;
QY 22 ILEVPESV---TGPWKGVNLPCTYDPLQGYTQVLVWLVQSGDPVTIFLRDSSGDHIQ 78
DB 143 ILEIPNSKLEDPQGGYSGHIENDSGDNCHGATILEPPYFVTPLPEPVQVTVGDSASLQCC 202
QY 79 QAK-----YQG-----RLVSHKVPQGVDSLSQLSTLEMDRSHYTCVWTQTP 120
DB 203 VAGTPEMIVSYKGDGTLRGATATVMHFKNQV---ATLVFQVDSDDSGEVICKVENTVG 259
QY 121 DGNQ-----VVRDKITELRVQKLSVKPTVTGSGYGTVPQGMRIISLQCGARGSPPIYS 175
DB 260 EATSSLLTVQERKLPSPFTRKLRDVHETV-----GLPVTDCGIAGEPIEV 307
QY 176 IWKYQQTNNQEPKIVAT---LSTLLFKPAVIADS---GSYFCTAKGVSEQHSIDIVKVFV 230
DB 308 SWFKDNVRVKEDYNVHTSFIDNVAAILQLTKSLMGQYTCYTAGNAIGT-----A 357
QY 231 KDSKSLKTKTEAPTTMTYPLKATSTVKQSWDVTDMDGVLGETS 275
DB 358 SSSGKLVLTGKTPPPFDTPI-----TPVDGIIGESA 389

RESULT 7

T42633
connectin/titin - chicken (fragment)
C:Species: Gallus gallus (chicken)
C>Date: 11-Jan-2000 #sequence_revision 11-Jan-2000 #text_change 09-Jul-2004
C:Accession: T42633
R:Yajima, H.; Ohtsuka, H.; Kawamura, Y.; Kume, H.; Murayama, T.; Abe, H.; Kimura, S.; Biochem. Biophys. Res. Commun. 223, 160-164, 1996
A:Title: A 11.5-kb 5'-terminal cDNA sequence of chicken breast muscle connectin/titin re
A:Reference number: 222221; MUID:96254045; PMID:8660363
A:Accession: T42633
A:Status: preliminary; translated from GB/EMBL/DBJ
A:Molecule type: mRNA
A:Residues: 1-4162 <YAJ>
A:Cross-references: UNIPROT:Q98918; EMBL:D83390; NID:G1513029; PIDN:BAAL1908.1; PID:G1513029
A:Experimental source: breast muscle
C:Keywords: skeletal muscle
Query Match 8.2%; Score 138.5; DB 2; Length 4162;
Best Local Similarity 21.4%; Pred. No. 0.076;
Matches 61; Conservative 38; Mismatches 117; Indels 69; Gaps 10;
QY 22 ILEVPESV---TGPWKGVNLPCTYDPLQGYTQVLVWLVQSGDPVTIFLRDSSGDHIQ 78

Db 3895 ILEIPNSKLEDOGOYSCHIENDSGQDNCHGAITILEPPYFVTPLEBPVQVTVGDSASLQCC 3954

Qy 79 QAK-----YQG-----RLVSHKVPDVSLSQLSTLEMDRSHYTCVTTWQTP 120

Db 3955 VAGTPEMIVSWYKGTDLRGATVAKHFNQV---ATLVFSQVSDSDSGEYICKVENTVG 4011

Qy 121 DGNQ-----VVRDKITELRVOKLSVSKPTVTGSGYGTVPQGMRIISLQCCQARGSPPIISY 175

Db 4012 EATSSLLTVQERKLPSPSTRKLRDVHETV-----GLPVTDCGIAGSEPIEV 4059

Qy 176 IWYKQQTNNQEPKIVAT--LSTLLFKPAVIADSGSYFCTAKQGVGEQHSIDIVKFWV 230

Db 4060 SFWKDNVRKEDYNNVHTSFIDNVAIQLIKTDKSLMGVYCTCTASNAIGT-----A 4109

Qy 231 KDSSKLLKTKTAPRTMYPLKATSTVKQSWDWTMDGILGETS 275

Db 4110 SSSGKLVLTGKTPPPFPDTP-----TPVDGIIGESA 4141

RESULT 8

T08678

hypothetical protein DKFZp564I11922.1 - human (fragment)

C:Species: Homo sapiens (man)

C>Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change 09-Jul-2004

C:Accession: T08678

R:Wambutt, R.; Heubner, D.; Mewes, H.W.; Gassenhuber, J.; Wiemann, S.

submitted to the Protein Sequence Database, May 1999

A:Reference number: Z16469

A:Accession: T08678

A:Molecule type: mRNA

A:Residues: 1-584 <WAM>

A:Cross-references: UNIPROT:O9Y3Y8; EMBL:AL049946

A:Experimental source: fetal brain; clone DKFZp564I11922

C:Genetics:

A:Note: DKFZp564I11922.1

Query Match 8.1%; Score 136; DB 2; Length 584;

Best Local Similarity 22.4%; Pred. No. 0.0099;

Matches 53; Conservative 37; Mismatches 99; Indels 48; Gaps 8;

Qy 35 GDNVLPCTVDPLQGYTVLWKLVQRGSDPVTIFLRDSSGDHIQOAKYQGRLHVSHKVP 94

Db 15 GDLKVDVCV---ATGLPNPEISMSLPDGSIVNSFMQSDSDSGRTKR-----YVFN 61

Qy 95 DVSLSQLSTLEMDRSHYTCVTTWQTPDGNQVVRDKITELRVOKLSVSKP-TVTTGSGYGF 153

Db 62 NGTLFNEVGMREEGDYTCFAE-----NQVGKD---EMRVRVKVTATPATIRNKTCLAV 112

Qy 154 TVPQGMRIISLQCCQARGSPPIISYIWY---KQQTNNQEPKIVATLSTLLFKPAVIADSGSY 209

Db 113 QVPYGDVVTVACEAKGEMPKTWLSPTNKV IPTSSEKVIQYQDGTLLIQAKQRSDSGNY 172

Qy 210 FCTAKQGVGSEQH-----SDIVKPVVNDSSKLLKTKTEA-PT 245

Db 173 TCLVRNSAGEDRKTVIHVNVQPPKINGPNPITTVREIAAGSGRKLIDCKRAEGIPT 229

RESULT 9

T20992

hypothetical protein F15G9.4a - Caenorhabditis elegans

C:Species: Caenorhabditis elegans

C>Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004

C:Accession: T20992; T24733

R:Sulston, J.

submitted to the EMBL Data Library, December 1994

A:Reference number: Z19355

A:Accession: T20992

A:Molecule type: DNA

A:Residues: 1-5175 <WIL>

A:Cross-references: UNIPROT:0810J3; EMBL:Z47068; PIDN:CAA87335.1; GSPDB:GN000028; CESP:F15G9

A:Experimental source: clone F15G9

R:Kershaw, J.

submitted to the EMBL Data Library, December 1994

A:Reference number: Z19929

A:Accession: T24733

A>Status: preliminary; translated from GB/EMBL/DBDJ

A:Molecule type: DNA

A:Residues: 1-5175 <WI2>

A:Cross-references: EMBL:Z47070; PIDN:CAA87344.1; GSPDB:GN000028; CESP:F15G9.4a

A:Experimental source: clone T09B9

C:Genetics:

A:Gene: CESP:F15G9.4a

A:Map position: X

A:Introns: 85/1; 334/3; 370/1; 477/2; 606/3; 664/1; 935/3; 977/1; 1051/3; 1184/3; 1201/1; 2512/2; 2593/3; 2699/3; 2759/1; 2852/1; 2889/3; 2913/3; 2941/1; 2967/3; 2991/3; 3033/1; 4225/1; 4361/1; 4408/1; 4456/1; 4498/1; 4647/3; 4838/1; 4879/1; 4941/1; 5011/1; 5077/1;

Query Match 7.9%; Score 133.5; DB 2; Length 5175;

Best Local Similarity 22.9%; Pred. No. 0.24;

Matches 48; Conservative 36; Mismatches 93; Indels 33; Gaps 6;

Qy 76 HIQAKYQGRLHVSHKVPDVSLSQLSTLEMDRSHYTCVTTWQTPDGNQVVRDKITELRV 135

Db 2796 HAHDESQVQGVITTSKYAANEKTLNVTNIQLDDEGFIYC-----TAVNEAGITKKFFKLIV 2850

Qy 136 -----QKLSVSKPTVTGSGYGTVPQGMRIISLQCCQARGSPPIISYIWYKQQTNNQE 186

Db 2851 IETPYFLDQOKL-----YPIILGKELTLDCSATGTPPTILPMKDGKRLNE 2896

Qy 187 PIKVATL-STLLFKPAVIADSGSYFCTAKQGVGEQHSIDIVKVV--KDSKLLKTKTEA 243

Db 2897 SDEVDIIIGSTLVIDNPQKEVEGRYTCTAENKAGRSEKDMVMVEVLLPPLKSKEMINVEQA 2956

Qy 244 --PTTMYPLKATSTVKQSWDWTMDGYL 271

Db 2957 GDPLTECPIDETSGVHITWSRQFGKDGQL 2986

RESULT 10

T43290

hemiceitin precursor - Caenorhabditis elegans

C:Species: Caenorhabditis elegans

C>Date: 11-Jan-2000 #sequence_revision 11-Jan-2000 #text_change 09-Jul-2004

C:Accession: T43290; T20993; T24734

R:Vogel, B.E.; Hedgecock, E.M.

submitted to the EMBL Data Library, June 1998

A>Description: Hemiceitin is required for hemidesmosome mediated cell adhesion and germ-

A:Reference number: Z22396

A:Accession: T43290

A>Status: preliminary; translated from GB/EMBL/DBDJ

A:Molecule type: mRNA

A:Residues: 1-5198 <VOG>

A:Cross-references: UNIPROT:O76518; EMBL:AF074901; PIDN:AAC26792.1

R:Sulston, J.

submitted to the EMBL Data Library, December 1994

A:Reference number: Z19355

A:Accession: T20993

A>Status: preliminary; translated from GB/EMBL/DBDJ

A:Molecule type: DNA

A:Residues: 1-5198 <WIL>

A:Cross-references: EMBL:Z47068; PIDN:CAA87336.1; GSPDB:GN000028; CESP:F15G9.4b

A:Experimental source: clone F15G9

R:Kershaw, J.

submitted to the EMBL Data Library, December 1994

A:Reference number: Z19929

A:Accession: T24734

A>Status: preliminary; translated from GB/EMBL/DBDJ

A:Molecule type: DNA

A:Residues: 1-5198 <WI2>

A:Cross-references: EMBL:Z47070; PIDN:CAA87345.1; GSPDB:GN000028; CESP:F15G9.4b

A:Experimental source: clone T09B9

C:Genetics:

A:Gene: him-4; F15G9.4b

A:Map position: X

A:Introns: 85/1; 120/1; 334/3; 370/1; 477/2; 606/3; 664/1; 935/3; 977/1; 1051/3; 1184/3;

2512/2; 2593/3; 2699/3; 2759/1; 2852/1; 2889/3; 2913/3; 2941/1; 2967/3; 2991/3; 3033/1
1; 4225/1; 4361/1; 4408/1; 4456/1; 4498/1; 4647/3; 4838/1; 4902/1; 4964/1; 5034/1; 5100/1

Query Match 7.9%; Score 133.5; DB 2; Length 5198;
Best Local Similarity 22.9%; Pred. No. 0.24;
Matches 48; Conservative 36; Mismatches 93; Indels 33; Gaps 6;

QY 76 HQQAQYQGRHLVSHKVPDVSQQLSTLEMDRSHYTCVTTWTPDGNQVVRDKITELRV 135
DB HAHDSEVQNGVITSKYAANEKTLNVTNIQLDDSGFYFC-----TAVNEAGITKKFFKLIV 2850

QY 136 -----QKLSVSKPTVTTCGSGYFTVQGNRISLQCAQSGSPISVIWYKQNTNOE 186
DB IETPYFLDQKQ-----YPIILGKRLTLDCASTGTPPTILFMKQGRKRLNE 2896

QY 187 PIKVAATL-STLLFKPAVIADSGSYFCTAKQGVGSEQHSIDIVKPVV--KDSSKLLTKTKEA 243
DB SDEVDIIGSTLVIDNPQKEVGYRTYCIENKAGRSEKMMVEVLLPPLKSKEMINVEVQA 2956

QY 244 --PTTWYPLKATSVKQSWDWTMDGYL 271
DB GDLPLTECIPTDTSVGHITWSRQFGKDGQL 2986

RESULT 11

PFHUGB
N:Contains: protein-tyrosine kinase (EC 2.7.1.112)
C:Species: Homo sapiens (man)
C:Date: 31-Dec-1992 #sequence_revision 31-Dec-1992 #text change 09-Jul-2004
C:Accession: A28206; A31195; A38268; A31925; B31925; C31925
R:Gronwald, R.G.K.; Grant, F.J.; Haldeman, B.A.; Hart, C.E.; O'Hara, P.J.; Hagen, F.S.;
Proc. Natl. Acad. Sci. U.S.A. 85, 3435-3439, 1988
A:Title: Cloning and expression of a cDNA coding for the human platelet-derived growth factor
A:Reference number: A28206; MUID:88217915; PMID:2835772
A:Accession: A28206
A:Molecule type: mRNA
A:Residues: 1-1106 <GRO>
A:Cross-references: UNIPROT:P09619; GB:J03278; NID:g189731; PIDN:AAA60049.1; PID:g189732
R:Clackson-Welsh, L.; Eriksson, A.; Moren, A.; Severinsson, L.; Ek, B.; Oestman, A.; Bet
Mol. Cell. Biol. 8, 3476-3486, 1988
A:Title: cDNA cloning and expression of a human platelet-derived growth factor (PDGF) re
A:Reference number: A31195; MUID:89096941; PMID:2850496
A:Accession: A31195
A:Molecule type: mRNA
A:Residues: 1-240.'D'.242-1106 <CLA>
A:Cross-references: GB:M21616; NID:g189729; PIDN:AAA36427.1; PID:g189730
R:Partanen, J.; Mäkelä, T.P.; Alitalo, R.; Lehtvaeslaihio, H.; Alitalo, K.
Proc. Natl. Acad. Sci. U.S.A. 87, 8913-8917, 1990
A:Title: Putative tyrosine kinases expressed in K-562 human leukemia cells.
A:Reference number: A38268; MUID:91062389; PMID:2247464
A:Accession: A38268
A>Status: nucleic acid sequence not shown
A:Molecule type: mRNA
A:Residues: 828-884 <PAR>
R:Roberts, W.M.; Look, A.T.; Rousset, M.F.; Sherr, C.J.
Cell 55, 655-661, 1988
A:Title: Tandem linkage of human CSF-1 receptor (c-fms) and PDGF receptor genes.
A:Reference number: A90908; MUID:89028677; PMID:2846185
A:Accession: A31925
A>Status: not compared with conceptual translation
A:Molecule type: DNA
A:Residues: 676-727 <ROB>
A:Accession: B31925
A>Status: not compared with conceptual translation
A:Molecule type: DNA
A:Residues: 901-932 <R02>
A:Accession: C31925
A>Status: not compared with conceptual translation
A:Molecule type: DNA
A:Residues: 1047-1106 <R03>
C:Comment: The extracellular domain is predicted to include five immunoglobulin-like dom
C:Genetics:

A:Gene: GDB:PDGFRB
A:Cross-references: GDB:120710; OMIM:173410
A:Map position: 5q31-5q32
C:Superfamily: macrophage colony-stimulating factor 1 receptor; immunoglobulin homology,
C:Keywords: ATP; autophosphorylation; glycoprotein; growth factor receptor; heterodimer,
F:1-32/Domain: signal sequence #status predicted <SIG>
F:33-1106/Product: platelet-derived growth factor receptor beta #status predicted <EXT>
F:33-531/Domain: extracellular #status predicted <IMM1>
F:47-102/Domain: immunoglobulin homology <IMM1>
F:142-192/Domain: immunoglobulin homology <IMM2>
F:228-293/Domain: immunoglobulin homology <IMM3>
F:429-510/Domain: immunoglobulin homology <IMM4>
F:532-555/Domain: transmembrane #status predicted <TM>
F:556-1106/Domain: intracellular #status predicted <INT>
F:598-965/Domain: protein kinase homology <KIN>
F:606-614/Region: protein kinase ATP-binding motif
F:45-89.103.215.230.292.307.354.371.468.479/Binding site: carbohydrate (Asn) (covalent)
F:54-100.149-190.235-291.436-508/Disulfide bonds: #status predicted
F:834/Active site: Lys #status predicted
F:857/Binding site: phosphate (Tyr) (covalent) (by autophosphorylation) #status predicted

Query Match 7.9%; Score 133; DB 1; Length 1106;
Best Local Similarity 21.4%; Pred. No. 0.038;
Matches 60; Conservative 41; Mismatches 92; Indels 88; Gaps 11;

QY 23 LEVPESVTGPW-----KGDVNLPTCTYDPLQYTVLVKLVQKSGDPVTIFLRDSSG 74
DB ITPICRVTDPLQVLTLLHEKKGDVALPVPYDHQGFSGI----- 182

QY 75 DHQQAQYQGRHLVSHKVPDVSQQLSTLEMDRSHYTCVTTWTPDGNQVVRDKITELRV 134
DB ITPICRVTDPLQVLTLLHEKKGDVALPVPYDHQGFSGI----- 182

QY 135 VQKLSVSKPTVTTCGSGYFTVQGNRISLQCAQSGSPISVIWYKQNTNO--EPIKVA 191
DB ITPICRVTDPLQVLTLLHEKKGDVALPVPYDHQGFSGI----- 182

QY 211 VSSINVSNAVQT-----VVRQGENITLMCIIVGNEVNVFWPTPRKESGRLEVPVTFD 264

QY 192 TL-----STLLFKPAVIADSGSYFCTAKQGVGSEQHSIDIVKPVVQKSSKLLTKTKEA 244
DB ITPICRVTDPLQVLTLLHEKKGDVALPVPYDHQGFSGI----- 182

QY 245 TMTVPLKATSVKQSWD-----WTTMDGYLGETSAG 277
DB ITPICRVTDPLQVLTLLHEKKGDVALPVPYDHQGFSGI----- 182

RESULT 12

C42632
cell adhesion molecule apCAM (clone d12) - California sea hare
C:Species: Aplysia californica (California sea hare)
C:Date: 04-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C:Accession: C42632
R:Mayford, M.; Barzilai, A.; Keller, F.; Schacher, S.; Kandel, E.R.
Science 256, 638-644, 1992
A:Title: Modulation of an NCAM-related adhesion molecule with long-term synaptic plasti
A:Reference number: A42632; MUID:92263095; PMID:1585176
A:Accession: C42632
A>Status: preliminary; not compared with conceptual translation
A:Molecule type: nucleic acid
A:Residues: 1-765 <NAY>
A:Cross-references: UNIPROT:Q9TWA4
A:Experimental source: CNS
A>Note: sequence extracted from NCBI backbone (NCBIP:101351)
C:Superfamily: neural cell adhesion molecule; fibronectin type III repeat homology; imm

Query Match 7.8%; Score 132.5; DB 2; Length 765;
Best Local Similarity 20.7%; Pred. No. 0.026;
Matches 54; Conservative 31; Mismatches 63; Indels 113; Gaps 10;

QY 104 EMDDRSHTYTCV-----TWQTPDGNQVVRDKITELRVQKLSVSKPT-----VTTGSG 150
DB ITPICRVTDPLQVLTLLHEKKGDVALPVPYDHQGFSGI----- 182

QY 230 KVGDEVKITCOATGVPPPTYPQFKGDVMTDEMWNNGVLTINPLKTTDQATYTCIATNKG 289

QY 151 YGFT-----VP-----QGMRIISLQCOARGSPPISYIW----- 177
|||
Db 290 -GFAESSNTLDVKVPPTIEDMEETDVAVSGQELTITCTAKGDPPEPSVIWKDGPQASD 348
|||
QY 178 -----YKQOTNNOEPIKVATLST-LLFKPAVIADSGSYFCTAKGQVGSQHSQSDIVKF 228
|||
Db 349 GIVNKGPTYEKVGSNQNDMEETVAQHMTFKPVTYQDAGTYICTAFSLVGSANKT--VKL 406
|||
QY 229 VVK-----DSSKLLKTKT 241
|||
Db 407 TVQYKPNFTDFKEREFFGWRGHKANLTCQANANPVATIEWMPDAENPDDYSKAVRIPN 466
|||
QY 242 EAPTMTYPLKATSTVKQSWD 262
|||
Db 467 EAPYTI-----NMLQKWD 479
|||
RESULT 13
B42632
cell adhesion molecule apCAM (clone d15) - California sea hare
C:Species: Aplysia californica (California sea hare)
C:Date: 04-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C:Accession: B42632
R:Mayford, M.; Barzilai, A.; Keller, F.; Schacher, S.; Kandel, E.R.
Science 256, 638-644, 1992
A:Title: Modulation of an NCAM-related adhesion molecule with long-term synaptic plastic
A:Reference number: A42632; MUID:92263095; PMID:1585176
A:Accession: B42632
A>Status: preliminary; not compared with conceptual translation
A:Molecule type: nucleic acid
A:Residues: 1-812 <MAY>
A:CROSS-references: UNIPROT:Q9TWA5
A:Experimental source: CNS
A>Note: sequence extracted from NCBI backbone (NCBIP:101346)
C:Superfamily: neural cell adhesion molecule; fibronectin type III repeat homology; immu

Query Match 7.8%; Score 132.5; DB 2; Length 812;
Best Local Similarity 20.7%; Pred. No. 0.028;
Matches 54; Conservative 31; Mismatches 63; Indels 113; Gaps 10;

QY 104 EMDDRSHTYTCV-----TWQTPDGNQVVRDKITELRVQKLSVSKPT-----VTGSG 150
|||
Db 230 KVGDEVKITCQATGVPPPTYQFKGDMVTDEMVGNGVLTINPLKTTDQATVTCIATNKG 289
|||
QY 151 YGFT-----VP-----QGMRIISLQCOARGSPPISYIW----- 177
|||
Db 290 -GFAESSNTLDVKVPPTIEDMEETDVAVSGQELTITCTAKGDPPEPSVIWKDGPQASD 348
|||
QY 178 -----YKQOTNNOEPIKVATLST-LLFKPAVIADSGSYFCTAKGQVGSQHSQSDIVKF 228
|||
Db 349 GIVNKGPTYEKVGSNQNDMEETVAQHMTFKPVTYQDAGTYICTAFSLVGSANKT--VKL 406
|||
QY 229 VVK-----DSSKLLKTKT 241
|||
Db 407 TVQYKPNFTDFKEREFFGWRGHKANLTCQANANPVATIEWMPDAENPDDYSKAVRIPN 466
|||
QY 242 EAPTMTYPLKATSTVKQSWD 262
|||
Db 467 EAPYTI-----NMLQKWD 479
|||
RESULT 14
A42632
cell adhesion molecule apCAM (clone d19) - California sea hare
C:Species: Aplysia californica (California sea hare)
C:Date: 04-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C:Accession: A42632
R:Mayford, M.; Barzilai, A.; Keller, F.; Schacher, S.; Kandel, E.R.
Science 256, 638-644, 1992
A:Title: Modulation of an NCAM-related adhesion molecule with long-term synaptic plastic
A:Reference number: A42632; MUID:92263095; PMID:1585176
A:Accession: A42632
A>Status: preliminary; not compared with conceptual translation

A:Molecule type: nucleic acid
A:Residues: 1-932 <MAY>
A:CROSS-references: UNIPROT:Q9TWA6
A:Experimental source: CNS
A>Note: sequence extracted from NCBI backbone (NCBIP:101342)
C:Superfamily: neural cell adhesion molecule; fibronectin type III repeat homology; immu

Query Match 7.8%; Score 132.5; DB 2; Length 932;
Best Local Similarity 20.7%; Pred. No. 0.033;
Matches 54; Conservative 31; Mismatches 63; Indels 113; Gaps 10;

QY 104 EMDDRSHTYTCV-----TWQTPDGNQVVRDKITELRVQKLSVSKPT-----VTGSG 150
|||
Db 230 KVGDEVKITCQATGVPPPTYQFKGDMVTDEMVGNGVLTINPLKTTDQATVTCIATNKG 289
|||
QY 151 YGFT-----VP-----QGMRIISLQCOARGSPPISYIW----- 177
|||
Db 290 -GFAESSNTLDVKVPPTIEDMEETDVAVSGQELTITCTAKGDPPEPSVIWKDGPQASD 348
|||
QY 178 -----YKQOTNNOEPIKVATLST-LLFKPAVIADSGSYFCTAKGQVGSQHSQSDIVKF 228
|||
Db 349 GIVNKGPTYEKVGSNQNDMEETVAQHMTFKPVTYQDAGTYICTAFSLVGSANKT--VKL 406
|||
QY 229 VVK-----DSSKLLKTKT 241
|||
Db 407 TVQYKPNFTDFKEREFFGWRGHKANLTCQANANPVATIEWMPDAENPDDYSKAVRIPN 466
|||
QY 242 EAPTMTYPLKATSTVKQSWD 262
|||
Db 467 EAPYTI-----NMLQKWD 479
|||
RESULT 15
S26180
neurofascin - chicken
C:Species: Gallus gallus (chicken)
C:Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 09-Jul-2004
C:Accession: S26180
R:Volkmert, H.; Hassel, B.; Wolff, J.M.; Frank, R.; Rathjen, F.G.
J. Cell Biol. 118, 149-161, 1992
A:Title: Structure of the axonal surface recognition molecule neurofascin and its relat
A:Reference number: S26180; MUID:92317154; PMID:1377696
A:Accession: S26180
A>Status: preliminary
A:Molecule type: mRNA
A:Residues: 1-1272 <VOL>
A:CROSS-references: UNIPROT:O42414; EMBL:X65224; NID:G63659; PIDN:CAA46330.1; PID:G63660
C:Superfamily: neural cell adhesion molecule I1; fibronectin type III repeat homology; i
F:279-336/Domain: immunoglobulin homology <IMM>

Query Match 7.8%; Score 132.5; DB 2; Length 1272;
Best Local Similarity 22.2%; Pred. No. 0.049;
Matches 51; Conservative 38; Mismatches 88; Indels 53; Gaps 7;

QY 22 ILEVPESVTGPMKGDVNLPCYDPLQGYTVLKVMLVORGSDPVTIFLDRSSGDHITQQA 81
|||
Db 150 VIEVDEG-----APLSLQCNPFP-----GLPPPVIFWSSSNEPIHQDK 188
|||
QY 82 -----YQGRLVHSHKVPGDVSLQLS-----TLEMDRSHYTCVETWQTPDGNQVVRDK 129
|||
Db 189 RVSGONGDLYFSNVMLQDAQTDYSCNARFHTHTTIQKNPYTLKVKTKKPHNETSLRNH 248
|||
QY 130 ITELAVQKLSVSKPT--VTGSGYGTVPQGNRIISLQCOARGSPPISYIWYK----- 179
|||
Db 249 TDMYSGRGVTEPTPFPYGYSSQVNLRGVLDLLECIASGVPAFDINWYKKGGLPAG 308
|||
QY 180 --QOTNNOEPIKVATLSTLLFKPAVIADSGSYFCTAKGQVGSQHSQSDIVK 227
|||
Db 309 KTKLENFNKALRISNVSE-----EDSGEYFCLASNKMGSIIRHTISVR 350
|||
Search completed: August 26, 2005, 19:39:58
Job time : 72 secs

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Db 181 QTNQPEIKVATLSTLLFKPAVIADSGSYFCTAKQGVSEHQSDIVKFWKDSKLLKTK 240
QY 241 TEAPTTMTYPLKATSTVKQSDWTTMDGYLGETSAGPKSLPVPFAILLIISLCMVVFT 300
Db 241 TEAPTTMTYPLKATSTVKQSDWTTMDGYLGETSAGPKSLPVPFAILLIISLCMVVFT 300
QY 301 MAYIMLCRKTSQOEHVYEAAAR 321
Db 301 MAYIMLCRKTSQOEHVYEAAAR 321

RESULT 2

QY279 ID Q9Y279 PRELIMINARY; PRT; 399 AA.
AC Q9Y279
DT 01-NOV-1999 (TRENBLrel. 12, Created)
DT 01-NOV-1999 (TRENBLrel. 12, Last sequence update)
DT 25-OCT-2004 (TRENBLrel. 28, Last annotation update)
DE 239Ig protein precursor (V-set and immunoglobulin domain containing 4).
DE Name=239Ig; Synonyms=VSI64;
GN Homo sapiens (Human).
OS Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RN [2]
RP SEQUENCE FROM N.A.
RX MEDLINE=20461865; PubMed=11004523; DOI=10.1016/S0167-4781(00)00131-7;
LA Langnese K., Colleaux L., Kloos D.U., Fontes M., Wiesacker P.;
RT "Cloning of 239Ig, a novel gene with immunoglobulin-like domains
RT located on human chromosome X";
RL Biochim. Biophys. Acta 1492:522-525(2000).
RN [2]

SEQUENCE FROM N.A.

RC TISSUE=Brain;
RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Ustin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Bosak S.A., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Kettman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smallos D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Marra M.A.;
RP "Generation and initial analysis of more than 15,000 full-length human
RP and mouse cDNA sequences";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN [3]

SEQUENCE FROM N.A.

RC TISSUE=Brain;
RX Director MGC Project;
RA Submitted (JUL-2001) to the EMBL/GenBank/DBJ databases.
RL EMBL; AJ132502; CAB51536.1; -.
DR EMBL; BC010525; AAH10525.1; -.
DR InterPro; IPR007110; IG-like.
DR InterPro; IPR003598; IG_c2.
DR Pfam; PF00047; ig; 1.
DR SMART; SM00408; IGC2; 1.
DR PROSITE; PS50835; IG_LIKE; 2.
KW Signal.
FT SIGNAL 1 19 Potential.
FT CHAIN 20 399 239Ig protein.
FT SEQUENCE 399 AA; 43987 MW; 735CA3BC58185035 CRC64;

Query Match 100.0%; Score 1688; DB 2; Length 399;
Best Local Similarity 100.0%; Pred. No. 2.2e-125;
Matches 321; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MGILLGLLLGLHLTVDTYGRPILEVPEVSVTPGKWDVNLCTYDPLQGYTVLVKWLVR 60
Db 1 MGILLGLLLGLHLTVDTYGRPILEVPEVSVTPGKWDVNLCTYDPLQGYTVLVKWLVR 60
QY 61 GSDPVTIFLRDSSGDHIQQAQYQGRHLVSHKVPDGVSLQSLTLEMDRSHYTCVTTWQTP 120
Db 61 GSDPVTIFLRDSSGDHIQQAQYQGRHLVSHKVPDGVSLQSLTLEMDRSHYTCVTTWQTP 120
QY 121 DGNQVVRDKITELRVKLSVSKPTVTGSGYGFVTPQGMRIISLQCAQSGSPPISYIWKQ 180
Db 121 DGNQVVRDKITELRVKLSVSKPTVTGSGYGFVTPQGMRIISLQCAQSGSPPISYIWKQ 180
QY 181 QTNQPEIKVATLSTLLFKPAVIADSGSYFCTAKQGVSEHQSDIVKFWKDSKLLKTK 240
Db 181 QTNQPEIKVATLSTLLFKPAVIADSGSYFCTAKQGVSEHQSDIVKFWKDSKLLKTK 240
QY 241 TEAPTTMTYPLKATSTVKQSDWTTMDGYLGETSAGPKSLPVPFAILLIISLCMVVFT 300
Db 241 TEAPTTMTYPLKATSTVKQSDWTTMDGYLGETSAGPKSLPVPFAILLIISLCMVVFT 300
QY 301 MAYIMLCRKTSQOEHVYEAAAR 321
Db 301 MAYIMLCRKTSQOEHVYEAAAR 321

RESULT 3

Q80WA3 ID Q80WA3 PRELIMINARY; PRT; 280 AA.
AC Q80WA3
DT 01-JUN-2003 (TRENBLrel. 24, Created)
DT 01-JUN-2003 (TRENBLrel. 24, Last sequence update)
DT 01-MAR-2004 (TRENBLrel. 26, Last annotation update)
DE Hypothetical protein BC025105.
GN Name=BC025105;
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
RN [1]
RN [2]
RP SEQUENCE FROM N.A.
RC STRAIN=FVB/N; TISSUE=Liver;
RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Ustin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Bosak S.A., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Kettman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smallos D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Marra M.A.;
RP "Generation and initial analysis of more than 15,000 full-length human
RP and mouse cDNA sequences";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN [2]
RN [3]
RP SEQUENCE FROM N.A.
RC STRAIN=FVB/N; TISSUE=Liver;
RX Strausberg R.;
RL Submitted (MAR-2002) to the EMBL/GenBank/DBJ databases.
RL EMBL; BC025105; AAH25105.1; -.
DR MGD; MGI:2679720; BC025105.

DR InterPro: IPR007110; Ig-like.
DR PROSITE: PS00835; IG_LIKE; 1.
KW Hypothetical protein.
SQ SEQUENCE 280 AA; 31467 MW; 3F099AB996CE8977 CRC64;
Query Match 43.1%; Score 728; DB 2; Length 280;
Best Local Similarity 47.0%; Pred. No. 1.6e-49;
Matches 151; Conservative 29; Mismatches 45; Indels 96; Gaps 2;
QY 1 MGILLGLLLGLHLYTDYGRPILEVPESVTGPWKGDVNLPTCTYDPLQGYTQVLVWKLVR 60
Db 1 MEISSGLFLGLHLVLTGYHPTLKPTPESTGTWKGDKVLCQIYDPLRGVQLVWKLVRH 60
QY 61 GSDPVTIFURDSGHHQOAKYQGRHLVSHKVPDGVSLQSLTLEMDRSHYTCVTTWPTP 120
Db 61 GSDSVTIFURDSTGHHQOAKYQGRHLVSHKVPDGVSLQSLTLEMDRSHYTCVTTWPTP 120
QY 121 DGNQVVRDKITELRVKLSVSKPTVTGSGYFTVPQGNRISLQCCARGSPPIISWYKQ 180
Db 121 DGNQVVRDKITELRVKLSVSKPTVTGSGYFTVPQGNRISLQCCARGSPPIISWYKQ 180
QY 181 QTNQEPKIKVATLSTLLPKPAVIADSGSYFCTAKQGVGSEHQSDIVKVFVKDSKLLTKK 240
Db 142 -----NPP-----RIN 144
QY 241 TEAPTTMTVPLKATSTVKQSDWTTDMOGLGETSAGPGKSLPVFAILLIISLCMWVPT 300
Db 145 TEAPTTLHSSLEATTIMSTSDLTNTGTGLEETIAGSRNLPFAIFIISLCCIVAVT 204
QY 301 MAYIMLCRTSQOEHVYEAR 321
Db 205 IPYILFRCTFOOEVYGVSR 225
RESULT 4
JAM1_HUMAN STANDARD; PRT; 299 AA.
AC Q9Y624;
DT 16-OCT-2001 (Rel. 40, Created)
DT 16-OCT-2001 (Rel. 40, Last sequence update)
DT 25-JAN-2005 (Rel. 46, Last annotation update)
DE Functional adhesion molecule 1 precursor (JAM) (Platelet adhesion
molecule 1) (FAM-1) (Platelet F11 receptor) (UNQ264/PRO301).
GN Name=F11r; Synonyms=JAM1, JCAM;
OS Homo sapiens (Human)
OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=99323940; PubMed=10395639;
RA Ozaki H., Ishii K., Horiuchi H., Arai H., Kawamoto T., Okawa K.,
RA Iwanatsu A., Kita T.;
RT "Combined treatment of TNF-alpha and IFN-gamma causes redistribution
of functional adhesion molecule in human endothelial cells.";
RL J. Immunol. 163:553-557(1999).
RN [2]
RP SEQUENCE FROM N.A.
RA Sobocka M.B., Sobocki T., Rushbrook J.I., Banerjee P., Weiss C.,
RA Kornacki E.;
RT "Molecular cloning and sequencing of the cDNA of F11 receptor, a novel
Ig superfamily member from human platelets.";
RL Submitted (NOV-1999) to the EMBL/GenBank/DBJ databases.
RN [3]
RP SEQUENCE FROM N.A.
RA Naik U.P., Naik M.U., DeLeon P., Spathala J.;
RT "Cloning and characterization of FAM-1, a novel platelet adhesion
molecule involved in platelet activation.";
RL Submitted (JUL-1999) to the EMBL/GenBank/DBJ databases.
RN [4]
RP SEQUENCE FROM N.A.
RC TISSUE=Brain;
RX MEDLINE=21154917; PubMed=11230166; DOI=10.1101/gr.154701;

RA Wiemann S., Weil B., Wellenreuther R., Gassenhuber J., Glassl S.,
RA Anorge W., Boecher M., Bloecher H., Bauersachs S., Blum H.,
RA Lauber J., Duesterhoeft A., Beyer A., Koehrer K., Strack N.,
RA Meves H.-W., Ottenwaelder B., Obermaier B., Tampe J., Heubner D.,
RA Wambutt R., Korn B., Klein M., Poustka A.;
RT "Towards a catalog of human genes and proteins: sequencing and
analysis of 500 novel complete protein coding human cDNAs.";
RL Genome Res. 11:422-435(2001).
RN [5]
RP SEQUENCE FROM N.A.
RX MEDLINE=22887296; PubMed=12975309; DOI=10.1101/gr.1293003;
RA Clark H.F., Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J.,
RA Chen J., Chow B., Chui C., Crowley C., Currell B., Deuel B., Dowd P.,
RA Eaton D., Foster J., Grimaldi C., Gu Q., Haas P.E., Heldens S.,
RA Huang A., Kim H.S., Klimowski L., Jin Y., Johnson S., Lee J.,
RA Lewis L., Liao D., Mark M., Robbie E., Sanchez C., Schoenfeld J.,
RA Seehagiri S., Simmons L., Singh J., Smith V., Stinson J., Vagts A.,
RA Vandlen R., Watanabe C., Wieand D., Woods K., Xie M.-H., Yansura D.,
RA Yi S., Yu G., Yuan J., Zhang M., Zhang Z., Goddard A., Wood W.I.,
RA Godowski P., Gray A.;
RT "The secreted protein discovery initiative (SPDI), a large-scale
effort to identify novel human secreted and transmembrane proteins: a
bioinformatics assessment.";
RL Genome Res. 13:2265-2270(2003).
RN [6]
RP SEQUENCE FROM N.A.
RX TISSUE=Ovary;
RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Ustin T.B., Tothiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullighy S.J.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalón D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Kettman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smalios D.E.,
RA Schnurch A., Schein J.E., Jones S.J.M., Marra M.A.;
RT "Generation and initial analysis of more than 15,000 full-length human
and mouse cDNA sequences.";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN [7]
RP SEQUENCE OF 28-42.
RX PubMed=15340161; DOI=10.1110/ps.04682504;
RA Zhang Z., Henzel W.J.;
RT "Signal peptide prediction based on analysis of experimentally
verified cleavage sites.";
RL Protein Sci. 13:2819-2824(2004).
RN [8]
RP FUNCTION: Seems to plays a role in epithelial tight junction
formation. Appears early in primordial forms of cell junctions and
recruits PARD3. The association of the PARD6-PARD3 complex may
prevent the interaction of PARD3 with JAM1, thereby preventing
tight junction assembly (By similarity). Plays a role in
regulating monocyte transmigration involved in integrity of
epithelial barrier. Involved in platelet activation.
RN [9]
RP SUBUNIT: Interacts with the first PDZ domain of PARD3. The
association between PARD3 and PARD6B probably disrupts this
interaction (By similarity).
RN [10]
RP SUBCELLULAR LOCATION: Type I membrane protein (Potential).
RN [11]
RP TISSUE SPECIFICITY: Localized at tight junctions of both
epithelial and endothelial cells.
RN [12]
RP SIMILARITY: Belongs to the immunoglobulin-like V-type domains.
RN [13]
RP SIMILARITY: Contains 2 immunoglobulin-like V-type domains.
RN [14]
RP This SWISS-PROT entry is copyright. It is produced through a collaboration
between the Swiss Institute of Bioinformatics and the EMBL outstation -

CC the European Bioinformatics Institute. There are no restrictions on its
 CC use by non-profit institutions as long as its content is in no way
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 CC entities requires a license agreement (See <http://www.isb-sib.ch/announce/>
 CC or send an email to license@isb-sib.ch).
 CC -----

DR EMBL; AF111713; RAD42050.1; -
 DR EMBL; AF207907; AAF22829.1; -
 DR EMBL; AF172398; AAD48877.1; -
 DR EMBL; AL136649; CAB66584.1; -
 DR EMBL; AY358896; AAO89255.1; -
 DR EMBL; BC001533; AAH01533.1; -
 DR PIR; A59406; S56749.
 DR PDB; INHQ; X-ray; A/B=25-233.
 DR Genew; HGNC:14685; FLIR.
 DR MIM; 605721; -.

DR GO; GO:0005911; C:intercellular junction; TAS.
 DR GO; GO:0006954; P:inflammatory response; TAS.
 DR InterPro; IPR007110; Ig-like.
 DR Pfam; PF00047; Ig; 2.
 DR PROSITE; PS00835; IG_LIKE; 2.

KW 3D-structure; Direct protein sequencing; Glycoprotein;
 KW Immunoglobulin domain; Repeat; Signal; Tight junction; Transmembrane.
 FT SIGNAL 1 27

FT CHAIN 28 299 Junctional adhesion molecule 1.
 FT DOMAIN 28 238 Extracellular (Potential).
 FT TRANSMEM 239 259 Potential.
 FT DOMAIN 260 299 Cytoplasmic (Potential).
 FT DOMAIN 28 125 Ig-like V-type 1.
 FT DOMAIN 135 228 Ig-like V-type 2.
 FT DISULFID 50 109 Potential.
 FT DISULFID 133 212 Potential.
 FT CARBOHYD 185 185 N-linked (GlcNAc...) (Potential).
 SQ SEQUENCE 299 AA; 32593 MW; D95DE2FEA23D2851 CRC64;

Query Match 10.6%; Score 178.5; DB 1; Length 299;
 Best Local Similarity 25.5%; Pred. No. 6e-06;
 Matches 65; Conservative 36; Mismatches 93; Indels 51; Gaps 11;

QY 1 MGILGLLLGLHLLVDTYGRPILEVPESVTPGPKGVDNLPCYDPLQGYTVLVKLVQR 60
 DB 17 LAITLLCSALGVSIVHS-SEPEVRIPEN-----NPVKLSLAY---SGFSSPRVEW---- 62

QY 61 GSDPTIFLSDSGHIOAKQGRHLVSHK-----VPCDVSLSQLSTLEMDRSHYTCV 115
 DB 63 -----KPDGDTTLVVCNNKITASYEDRVTFLEPTGITFKSVTRE--DTGTYTCV 111

QY 116 TWQTPDGNQVVDKITELRVQKLSVSKPTVTGSGVGTVPQGMRIQLQCOAR-GSPPTIS 174
 DB 112 SEEGSGYGEVVKLVILV-----VPPSKPTVINPS-----SATIGNRAVLTCSEQDGSPPSE 163

QY 175 YTWYK-----QQTN-----NQEPKIVATLSTLFKPAVIADSGSYCTAKQGVGSQH 222
 DB 164 YTWFKDGIWMTNPKNKSTRAFSNSSLVNLPTTGELVFDPLSADTGEYSCEARNGYGTPT 223

QY 223 SDIVK 227
 DB 224 SNAVR 228

RESULT 5
 A33_HUMAN
 ID A33_HUMAN STANDARD; PRT; 319 AA.
 AC Q99795;
 DT 01-NOV-1997 (Rel. 35, Created)
 DT 01-NOV-1997 (Rel. 35, Last sequence update)
 DE 05-JUL-2004 (Rel. 44, Last annotation update)
 DE Cell surface A33 antigen precursor (Glycoprotein A33).
 GN Name=GPA33;
 OS Homo sapiens (Human).
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
 NCBI_TaxID=9606;

RN SEQUENCE FROM N.A., AND PARTIAL SEQUENCE.
 RP TISSUE=Colon carcinoma;
 RX MEDLINE=97165045; PubMed=9012807; DOI=10.1073/pnas.94.2.469;
 RA Heath J.K., White S.J., Johnstone C.N., Catmel B., Simpson R.J.,
 RA Moritz R.L., Tu G.-F., Ji H., Whitehead R.H., Groenen L.C.,
 RA Scott A.W., Ritter G., Cohen L., Welt S., Old L.J., Nice E.C.,
 RA Burgess A.W.;
 RT "The human A33 antigen is a transmembrane glycoprotein and a novel
 RT member of the immunoglobulin superfamily";
 RL Proc. Natl. Acad. Sci. U.S.A. 94:469-474(1997).
 RN [2]
 RP POST-TRANSLATIONAL MODIFICATIONS.
 RX MEDLINE=97396159; PubMed=9245713; DOI=10.1006/bbrc.1997.6966;
 RA Ritter G., Cohen L.S., Nice E.C., Catmel B., Burgess A.W.,
 RA Moritz R.L., Ji H., Heath J.K., White S.J., Welt S., Old L.J.,
 RA Simpson R.J.;
 RT "Characterization of posttranslational modifications of human A33
 RT antigen, a novel palmitoylated surface glycoprotein of human
 RT gastrointestinal epithelium";
 RL Biochem. Biophys. Res. Commun. 236:682-686(1997).
 CC -!- FUNCTION: May play a role in cell-cell recognition and signaling.
 CC -!- SUBCELLULAR LOCATION: Type I membrane protein.
 CC -!- TISSUE SPECIFICITY: Expressed in normal gastrointestinal
 CC epithelium and in 95% of colon cancers.
 CC -!- PTM: N-glycosylated, contains approximately 8 kDa of N-linked
 CC carbohydrate.
 CC -!- PTM: Palmitoylated.
 CC -!- SIMILARITY: Contains 1 immunoglobulin-like C2-type domain.
 CC -!- SIMILARITY: Contains 1 immunoglobulin-like V-type domain.
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 CC -----

DR EMBL; U79725; AAC50957.1; -.
 DR HSSP; O88792; 1F97.
 DR Genew; HGNC:4445; GPA33.
 DR MIM; 602171; -.

DR GO; GO:0005887; C:integral to plasma membrane; TAS.
 DR GO; GO:0004872; F:receptor activity; TAS.
 DR InterPro; IPR007110; Ig-like.
 DR Pfam; PF00047; Ig; 2.

DR PROSITE; PS00835; IG_LIKE; 2.
 DR Antigen; Direct protein sequencing; Glycoprotein;
 DR Immunoglobulin domain; Lipoprotein; Palmitate; Signal; Transmembrane.

FT SIGNAL 1 21
 FT CHAIN 22 319 Cell surface A33 antigen.
 FT DOMAIN 22 235 Extracellular (Potential).
 FT TRANSMEM 236 256 Potential.
 FT DOMAIN 257 319 Cytoplasmic (Potential).
 FT DOMAIN 22 134 Ig-like V-type.
 FT DOMAIN 140 227 Ig-like C2-type.
 FT DOMAIN 258 261 Poly-Cys.
 FT DISULFID 43 117 Potential.
 FT DISULFID 146 222 Potential.
 FT DISULFID 162 211 Potential.
 FT CARBOHYD 112 112 N-linked (GlcNAc...)
 FT CARBOHYD 200 200 N-linked (GlcNAc...)
 FT CARBOHYD 223 223 N-linked (GlcNAc...)
 SQ SEQUENCE 319 AA; 35632 MW; 9BFC7AAF45C2408E CRC64;

Query Match 10.5%; Score 177; DB 1; Length 319;
 Best Local Similarity 27.5%; Pred. No. 8.6e-06;
 Matches 64; Conservative 39; Mismatches 94; Indels 36; Gaps 11;

QY 4 LLGALLGLHLLVDTYGRPILEVPESVTPGPKGVDNLPCYDPLQGYTVLVKLVQR 60
 DB 8 VLWTLCAVRVTDAIS---VETPDVLRASQGSVTLPTCTVHTSTSSREGLIQWDKLLLT 64

```
QY 61 GSDPVTIFLRDSSGDHIOQAKYQGRHLVSHKV-PGDVSLQLSTLEMDRSHYTCVWTQT 119
Db 65 HTERVWVW-PFSKNYIHGELYKRVSVISNNAEQSDASITIDQLTMADNGTYECSVLSMS 123
QY 120 P-DGNQVRDKITELRVOKLSVSKPTVTGSGYGTVPQGMRIISLQCAR-GSPISVIW 177
Db 124 DLEGN-----TKGRVLLVLPVPSKPECGETI-IGNNIQTCSKESGTPQYSW 175
QY 178 YKQQTNNQ-----PIKVATLSTLLFKPAVIADSGSYFCTAKGQVQSE 220
Db 176 KRYNINLQEOPLAQPASQGVSLKNIST-----DTSGYYICTSSNEEGTQ 220

RESULT 6
Q10465 PRELIMINARY; PRT; 7962 AA.
AC Q10465;
DT 01-NOV-1996 (TREMBLrel. 01, Created)
DT 01-NOV-1996 (TREMBLrel. 01, Last sequence update)
DT 01-OCT-2003 (TREMBLrel. 25, Last annotation update)
DE Elastic titin (Fragment).
GN Name=titin;
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=Berkley; TISSUE=Skeletal muscle;
RX MEDLINE=96026330; PubMed=7569978;
RA Labelit S., Kolmerer B.;
RT "Titins, giant proteins in charge of muscle ultrastructure and
RT elasticity."
RL Science 270:293-296(1995).
DR EMBL; X90589; CA62189.1; -.
DR PIR; I38346; I38346.
DR HSSP; P11362; 1EVT.
DR GO; GO:0030017; C:sarcomere; TAS.
DR GO; GO:0008307; P:structural constituent of muscle; TAS.
DR GO; GO:0005941; P:striated muscle contraction; TAS.
DR InterPro; IPR007110; IG-like.
DR InterPro; IPR003598; IG_c2.
DR InterPro; IPR004168; PPAK_motif.
DR SMART; SM00408; IGC2; 43.
DR PROSITE; PS50835; IG_LIKE; 58.
FT NON_TER 1
FT NON_TER 7962
SQ SEQUENCE 7962 AA; 883018 MW; B85240533CBAD58 CRC64;

Query Match 10.2%; Score 172; DB 2; Length 7962;
Best Local Similarity 27.5%; Pred. No. 0.0013;
Matches 67; Conservative 30; Mismatches 85; Indels 62; Gaps 11;

QY 16 DTYGRPILEVPEPES-VTGPWKGDVNLNLP-----CTYDPLQGYTVQLVKW-----LVORGSDP 64
Db 860 DSSGALIVQEPSPFVTKPSKDV-LPGSAVCLKSTFGSTPLTIWFKGNKELVSGSCY 918
QY 65 VTIFLRDSSGDHIOQAKYQGRHLVSHKVPGDVSLQLSTLEMDRSHYTCVWTQPDG-- 122
Db 919 IT-----KEALESSELYLVKTSDSGTGYTCVKS-NVAGVGVE 953
QY 123 ---NQVRDKITELRVOKLSVSKPTVTGSGYGTVPQGMRIISLQCARGPSPIYVWK 179
Db 954 CSANLFWKEPAT--FVEKLEPQ-----LLKKGDTATQLACKVTGTPPIKITWPA 1000
QY 180 QQTNNQEPK-----VATLSTLLFKPAVIADSGSYFCTAKGQVQSEHQSHDIKVFVKDSS 234
Db 1001 NDREIKESKSRMSFVESTAVLRITDVGIEDSGEYWCENAGSDHCSIV--IVKESP 1058
QY 235 KULK 238
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Db 1059 YFTK 1062

RESULT 7
QBWZ42 PRELIMINARY; PRT; 34350 AA.
ID QBWZ42;
AC QBWZ42;
DT 01-MAR-2002 (TREMBLrel. 20, Created)
DT 01-MAR-2002 (TREMBLrel. 20, Last sequence update)
DT 01-MAR-2004 (TREMBLrel. 26, Last annotation update)
DE Titin.
GN Name=TTN;
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=20309627; PubMed=10850961;
RA Freiburg A., Trombitas K., Hell W., Cazorla O., Fougereousse F.,
RA Centner T., Kolmerer B., Witt C., Beckmann J.S., Gregorio C.C.,
RA Granzier H., Labelit S.;
RT "Series of exon-skipping events in the elastic spring region of titin
RT as the structural basis for myofibrillar elastic diversity."
RL Circ. Res. 86:1114-1121(2000).
RN [2]
RP SEQUENCE FROM N.A.
RX MEDLINE=21573839; PubMed=11717165;
RA Bang M.L., Centner T., Fornoff F., Geach A.J., Gotthardt M.,
RA Mcnabb M., Witt C.C., Labelit D., Gregorio C.C., Granzier H.,
RA Labelit S.;
RT "The complete gene sequence of titin, expression of an unusual ~700
RT kDa titin isoform and its interaction with obscurin identify a novel
RT Z-line to I-band linking system."
RL Circ. Res. 89:1065-1072(2001).
RN [3]
RP SEQUENCE FROM N.A.
RA Centner S.B.;
RL Submitted (JUL-2001) to the EMBL/GenBank/DBJ databases.
DR EMBL; AJ277892; CAD12456.1; -.
DR PDB; 1TNM; NMR; @=33480-33579.
DR PDB; 1TNN; NMR; @=33480-33579.
DR Genew; HGNC:12403; TTN.
DR GO; GO:0016020; C:membrane; IEA.
DR GO; GO:0005524; P:ATP binding; IEA.
DR GO; GO:0004896; P:hematopoietin/interferon-class (D200-domain. . .; IEA.
DR GO; GO:0004674; P:protein serine/threonine kinase activity; IEA.
DR GO; GO:0008307; P:structural constituent of muscle; IEA.
DR GO; GO:0007517; P:muscle development; IEA.
DR GO; GO:0006468; P:protein amino acid phosphorylation; IEA.
DR Pfam; PF00041; fn3; 132.
DR Pfam; PF00047; ig; 3.
DR Pfam; PF00069; Pkinase; 1.
DR Pfam; PF02818; PPAK; 53.
DR ProDom; PD000001; Prot_kinase; 1.
DR SMART; SM00060; FN3; 132.
DR SMART; SM00408; IGC2; 65.
DR SMART; SM00220; S_TKG; 1.
DR PROSITE; PS00933; FCGY_KINASES_1; UNKNOWN_1.
DR PROSITE; PS50853; FN3; 132.
DR PROSITE; PS50835; IG_LIKE; 143.
DR PROSITE; PS00290; IG_MHC; UNKNOWN_1.
DR PROSITE; PS00435; PEROXIDASE_1; UNKNOWN_1.
DR PROSITE; PS50011; PROTEIN_KINASE_DOM; 1.
DR PROSITE; PS50011; PROTEIN_KINASE_TYR; UNKNOWN_1.
DR PROSITE; PS00109; PROTEIN_KINASE_TYR; UNKNOWN_1.
SQ SEQUENCE 34350 AA; 3816189 MW; SB1120058A7CE58A CRC64;

Query Match 10.2%; Score 172; DB 2; Length 34350;
Best Local Similarity 27.5%; Pred. No. 0.0084;
Matches 67; Conservative 30; Mismatches 85; Indels 62; Gaps 11;

QY 16 DTYGRPILEVPEPES-VTGPWKGDVNLNLP-----CTYDPLQGYTVQLVKW-----LVORGSDP 64
Db 16 DTYGRPILEVPEPES-VTGPWKGDVNLNLP-----CTYDPLQGYTVQLVKW-----LVORGSDP 64
```

Db 5122 DSSGALIVQPPSFVTPKGSKDV-LPGSAVCLKSTFQGSTPLTIWRFGKNKELVSGGSCY 5180
QY 65 VTIFLRDSSGDHIQAKYQGRHLVSHKVPDVSLSQLEMDRSHYTCVETWQTPDG-- 122
Db 5181 IT-----KEALESLELYLVKTSDSGTYTCRVS-NVAGGVE 5215
QY 123 ---NOVVRDKITELRVQKLSVSKPTVTTGSGYGFVPOGMRISLQCAQSGPPISYIWK 179
Db 5216 CSANLFVKEPAT--FVEKLEPSQ-----LLKKGDATAQLACKVTGPPIKITWFA 5262
QY 180 QQTNNQEPK-----VATLSTLLFKPAVADSGSYFCTAKQGVGSEQHSIDIVKFVVKDSS 234
Db 5263 NDREIKESSKHRMSFVSTAVLRLLTDVGIEDSGEYCEAQAEGSDHCSSIV--IVKESP 5320
QY 235 KLLK 238
Db 5321 YFTK 5324

RESULT 8

Q66172 ID Q66172 PRELIMINARY; PRT; 292 AA.
AC Q66172;
DT 25-OCT-2004 (TReMBLrel. 28, Created)
DT 25-OCT-2004 (TReMBLrel. 28, Last sequence update)
DT 25-OCT-2004 (TReMBLrel. 28, Last annotation update)
DE Zgc:103642.
GN Name=zgc:103642;
OS Brachydanio rerio (Zebrafish) (Danio rerio).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;
OC Cyprinidae; Danio.
OX NCBI_TaxID=7955;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Larvae;
RX PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
PA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Bosak S.A., McKwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Kettaman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J.J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Marra M.A.;
RT "Generation and initial analysis of more than 15,000 full-length human
and mouse cDNA sequences.";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN [2]

SEQUENCE FROM N.A.

RC TISSUE=Larvae;
RA Director MGC Project;
RL Submitted (SEP-2004) to the EMBL/GenBank/DBSJ databases.
DR EMBL; BC081502; AH81502.1; -.
DR InterPro; IPR003599; IG.
DR InterPro; IPR007110; IG-like.
DR InterPro; IPR003598; IG_c2.
DR Pfam; PF00047; ig; 2.
DR SMART; SM00409; IG; 2.
DR SMART; SM00408; IGC2; 2.
DR PROSITE; PS50835; IG_LIKE; 2.
SQ SEQUENCE 292 AA; 32091 MW; E1F6B5FE74A72502 CRC64;

Query Match

10.1%; Score 170.5; DB 2; Length 292;

Best Local Similarity 28.6%; Pred. No. 2.5e-05;
Matches 64; Conservative 25; Mismatches 94; Indels 41; Gaps 10;
QY 28 SVTPGPK-----GDVNLCTYDPLQGYQVLVVKLVQ--RGSDPVTIFLRDSSGDHIQOAK 81
Db 22 TVTSPVKVKEGVDLQCSYTSDFGATP-RVEMFKDLKGSQTLVVF-----DCKPTGQ 74
QY 82 YQGRHLVSHKVPDVSLSQLEMDRSHYTCVETWQTPDGNOVVRDKITELRVQKLSVS 141
Db 75 YTGRTVMY-----DKLRFNFKVTRADTGDYDCEVSGSGGYGENTI-----KLTVL 119
QY 142 KPTVTGSGYGFVPOGMRISLQCAQSGPPISYIWKQQTN-NQBPPIKVATLSTLLFK 199
Db 120 VPPAKPVSRISSVTTSSNVRLTCFDPVGSPPSYIKYKONTPLPEDPTKPPAFKNLYTK 179
QY 200 -----PAVI-ADSGSYFCTAKQGVGSEQHSIDIVKFVVKD 232
Db 180 MNVFNGLFPSPSKMDTGSYFCEASNGEGVPQRGDEVKMEVRD 223

RESULT 9

JAMI BOVIN ID JAMI BOVIN STANDARD; PRT; 298 AA.
AC Q9XT56;
DT 16-OCT-2001 (Rel. 40, Created)
DT 16-OCT-2001 (Rel. 40, Last sequence update)
DT 05-JUL-2004 (Rel. 44, Last annotation update)
DE Junctional adhesion molecule 1 precursor (JAM).
GN Name=JAM1; Synonyms=JAM1;
OS Bos taurus (Bovine).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;
OC Bovinae; Bos.
OX NCBI_TaxID=9913;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=99323940; PubMed=10395639;
RA Ozaki H., Ishii K., Horiuchi H., Arai H., Kawamoto T., Okawa K.,
RA Iwamatsu A., Kita T.;
RT "Combined treatment of TNF-alpha and IFN-gamma causes redistribution
of junctional adhesion molecule in human endothelial cells.";
RL J. Immunol. 163:553-557(1999).
CC -!- FUNCTION: Seems to plays a role in epithelial tight junction
formation. Appears early in primordial forms of cell junctions and
recruits PAR3. The association of the PAR6-PAR3 complex may
prevent the interaction of PAR3 with JAM1, thereby preventing
tight junction assembly (By similarity). Plays a role in
regulating monocyte transmigration involved in integrity of
epithelial barrier. Involved in platelet activation.
CC -!- SUBUNIT: Interacts with the first PDZ domain of PAR3. The
association between PAR3 and PAR6B probably disrupts this
interaction (By similarity).
CC -!- SUBCELLULAR LOCATION: Type I membrane protein (Potential).
CC -!- TISSUE SPECIFICITY: Localized at tight junctions of both
epithelial and endothelial cells.
CC -!- SIMILARITY: Belongs to the immunoglobulin superfamily.
CC -!- SIMILARITY: Contains 2 immunoglobulin-like V-type domains.
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or send an email to license@isb-sib.ch).
CC -----
DR EMBL; AF111714; AAD42051.1; -.
DR HSSP; Q9Y624; 1NBQ.
DR InterPro; IPR007110; IG-like.
DR InterPro; IPR003598; IG_c2.
DR Pfam; PF00047; ig; 2.
DR SMART; SM00408; IGC2; 1.
DR PROSITE; PS50835; IG_LIKE; 2.

KW Glycoprotein; Immunoglobulin domain; Repeat; Signal; Tight junction;
KW Transmembrane.
FT SIGNAL 1 24 Potential.
FT CHAIN 25 237 Junctional adhesion molecule 1.
FT DOMAIN 25 298 Extracellular (Potential).
FT TRANSMEM 238 258 Potential.
FT DOMAIN 259 298 Cytoplasmic (Potential).
FT DOMAIN 28 124 Ig-like V-type 1.
FT DISULFID 134 227 Ig-like V-type 2.
FT DOMAIN 134 227 Potential.
FT DISULFID 49 108 Potential.
FT CARBOHYD 152 211 Potential.
FT CARBOHYD 184 184 N-linked (GlcNAc...) (Potential).
SQ SEQUENCE 298 AA; 32456 MW; 714FE1C1714769A2 CRC64;

Query Match 10.0%; Score 169.5; DB 1; Length 298;
Best Local Similarity 25.1%; Pred. No. 3.1e-05;
Matches 63; Conservative 31; Mismatches 88; Indels 69; Gaps 11;

QY 3 ILGLGLLGLHLYTDYGRILEVPESVTPGPKGDVNLFCYDPLQGYTQVLVQKVLVQRG- 61
DB 18 MILCSLALGRGAVQTY-EPVVRVPE-----NPAKLSCSY---SGFSRPMVKFTHGD 67
QY 62 -----SDPVTIFLRDSSGDHIQAKYQGRHLVSHKVPDGVSLQLSTLMDRSHYTC 114
DB 68 RGLVYCNKIT-----ASVNRVTF-----DTGITFHVTRKDTGMYTOM 109
QY 115 VTWQTPDGNQVVRDKITELRVOKLSVSKPTVTGSGYGTVPQGMRLSQCOAR-GSPPI 173
DB 110 V5-----DEGGNTYGEVTLVLPSPKPTINVP-----SVTIGTRAVLTCSRDSGSPS 161
QY 174 SYIWK-----OQTWNQEPKIVATLSTLLFKPAVIADSGSYFCTAKG 215
DB 162 EYKFKDGVEMPLEPKSNRAFSNSSTYLNQK-----TGELIFDPSASDTGDFTCQAQ 215
QY 216 QVGSQSHSDIV 226
DB 216 GYASPVKSDTV 226

RESULT 10
Q8WV3 PRELIMINARY; PRT; 365 AA.
ID Q8WV3
AC Q8WV3
DT 01-MAR-2002 (TrEMBLrel. 20, Created)
DT 01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Cocksackie virus and adenovirus receptor BCAR.
OS Bos taurus (Bovine).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;
OC Bovinae; Bos.
OX NCBI_TaxID=9913;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=21547769; PubMed=11689979; DOI=10.1006/bbrc.2001.5851;
RA Thoenen I., Keyaerts E., Lindberg M., Van Ranst M.;
RT "Characterization of a cDNA encoding the bovine coxsackie and
RT adenovirus receptor.";
RL Biochem. Biophys. Res. Commun. 288:805-808(2001).
DR EMBL; AY033651; AAKS7804.1; -.
DR FIR; JC7780; JC7780.
DR HSSP; P78310; IKAC.
DR GO; GO:0004872; F:receptor activity; IEA.
DR SMART; SM00408; IGC2; 1.
DR PROSITE; PS50835; IG_LIKE; 2.
KW Receptor.
SQ SEQUENCE 365 AA; 40153 MW; 36DE0BE5DCF86CF9 CRC64;

Query Match 10.0%; Score 169; DB 2; Length 365;
Best Local Similarity 24.8%; Pred. No. 4.4e-05;
Matches 85; Conservative 40; Mismatches 120; Indels 98; Gaps 17;

QY 1 MGILGLLLGLHLYTDYGRILEVPESVTPGPKGD-VNLPCTY-----DPLQGYTQV 52

DB 1 MELLRLFLLCGVADFTRLGSI-TTPEQMIEKAGETAYLFCFTLGPEDQGPLD----- 54
QY 53 LVKVLVQGRS-----DPVTIFLRDSSGDHIQAKYQ-----GRHLVSHK---VPGDVSLLQSTL 103
DB 55 -IEWLSPADNQKVDQVILLY---SGDKYDDYQDLKGRVHFTSNDLKSGDASINVTNL 110
QY 104 EMDRSHYTCVETWQTPDGNQVVRDKITELRVOKLSVSKPTVTGSGYGTVPQ-----GM 159
DB 111 QLSDIGTYQCKVKKAPGVGNK-----KIQLTVLWK-----SGIRCVDGSEIGN 156
QY 160 RISLQCOAR-GSPISYIWKYQQTNNQEP-----IKVATLSTLLFKPAVIADSGSYFCTAKG 215
DB 157 DFKLKCEPKGSLFLRYEWQKLSQKLPSTSWLPSTSPVISVKNASAEYSGTCTYVRN 216
QY 216 QVGSQSHSDIVKFFVVKDSSKLLTKTEATPTMTYPLKATSTVKQSDWTTDMDGYLGSTS 275
DB 217 RVGSDQ-----CLLRDVPFNSR-----AGTI 239
QY 276 AGPKSIPVFAILIILSCMVFTMAYIMLCRKTSQOEHVYE 318
DB 240 AG-----AVIGTLALVLIALIVE-----CCHKRREKYE 270

RESULT 11
Q8VC39 PRELIMINARY; PRT; 300 AA.
ID Q8VC39
AC Q8VC39
DT 01-MAR-2002 (TrEMBLrel. 20, Created)
DT 01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
DT 25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
DE Flii protein (Mus musculus adult male cecum cDNA, RIKEN full-length
DE enriched library, clone:9130004G24 product:junction cell adhesion
DE molecule1, full insert sequence).
GN Name=Flii;
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=mix FVB/N; TISSUE=Mammary tumor;
RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.2426038999;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Heieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Ustin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Kettman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Marra M.A.;
RT "Generation and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences.";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN [2]
RP SEQUENCE FROM N.A.
RC STRAIN=mix FVB/N; TISSUE=Mammary tumor;
RA Strausberg R.;
RN Submitted (JAN-2002) to the EMBL/GenBank/DBJ databases.
RN [3]
RP SEQUENCE FROM N.A.
RC STRAIN=CS7BL/6J; TISSUE=Cecum;
RX MEDLINE=99279253; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9;
RA Carninci P., Hayashizaki Y.;

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GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: August 26, 2005, 19:32:58 ; Search time 43 Seconds
(without alignments)
557.264 Million cell up

Title: US-10-767-374-2
 Perfect score: 1688
 Sequence: 1 MGILLGLLLGLHLTVDTYGR.....AYIMLCRTSQOEHHVEAAR 321

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 513545 seqs, 74649064 residues

Total number of hits satisfying chosen parameters: 513545

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Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                  Maximum Match 10%
                  Listing first 15

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3: /cgn2_6/ptodata/1/iaa/6A COMB. pep: *
4: /cgn2_6/ptodata/1/iaa/6B COMB. pep: *
5: /cgn2_6/ptodata/1/iaa/PCTUS COMB. pep: *
6: /cgn2_6/ptodata/1/iaa/backfiles1.per
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB	ID	Description
1	1688	100.0	321	4	US-09-254-465A-2	Sequence 2, Appli
2	1688	100.0	321	4	US-09-953-499-2	Sequence 2, Appli
3	1137	67.4	306	4	US-09-369-247-63	Sequence 63, Appl
4	870	51.5	175	4	US-09-763-902B-6	Sequence 6, Appli
5	178.5	10.6	299	3	US-09-188-930-189	Sequence 189, App
6	178.5	10.6	299	3	US-09-188-930-331	Sequence 331, App
7	178.5	10.6	299	3	US-09-462-270-2	Sequence 2, Appli
8	178.5	10.6	299	4	US-09-254-465A-1	Sequence 1, Appli
9	178.5	10.6	299	4	US-09-312-283C-189	Sequence 189, App
10	178.5	10.6	299	4	US-09-312-283C-331	Sequence 331, App
11	178.5	10.6	299	4	US-09-907-794A-119	Sequence 119, App
12	178.5	10.6	299	4	US-09-905-125A-119	Sequence 119, App
13	178.5	10.6	299	4	US-09-902-775A-119	Sequence 119, App
14	178.5	10.6	299	4	US-09-397-243D-3	Sequence 3, Appli
15	178.5	10.6	299	4	US-09-906-700-119	Sequence 119, App
16	178.5	10.6	299	4	US-09-903-603A-119	Sequence 119, App
17	178.5	10.6	299	4	US-09-904-920A-119	Sequence 119, App
18	178.5	10.6	299	4	US-09-909-064-119	Sequence 119, App
19	178.5	10.6	299	4	US-09-905-381A-119	Sequence 119, App
20	178.5	10.6	299	4	US-09-906-618-119	Sequence 119, App
21	178.5	10.6	299	4	US-09-953-499-1	Sequence 1, Appli
22	177	10.5	319	1	US-08-597-495B-22	Sequence 22, Appl
23	177	10.5	319	3	US-09-068-051A-22	Sequence 22, Appl
24	177	10.5	319	4	US-09-336-536-67	Sequence 67, Appl
25	177	10.5	319	4	US-09-254-465A-6	Sequence 6, Appli
26	177	10.5	319	4	US-09-953-499-6	Sequence 6, Appli
27	176	10.4	270	4	US-09-254-465A-24	Sequence 24, Appl


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Db 241 TEAPTTMTYPLKATSTVKQSDWTTMDGVLGETSAGPGKSLPVFAIILISLCCMWVFT 300
QY 301 MAYIMLCRKTSQOEHVYEAAAR 321
Db 301 MAYIMLCRKTSQOEHVYEAAAR 321

RESULT 2
US-09-953-499-2
; Sequence 2, Application US/09953499
; Patent No. 6838554
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Fong, Sherman
; APPLICANT: Goddard, Audrey
; APPLICANT: Gurney, Austin L.
; APPLICANT: Napier, Mary A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Wood, William I.
; TITLE OF INVENTION: COMPOSITIONS, COMPOSITIONS AND METHODS FOR THE TREATMENT
; OF DISEASES CHARACTERIZED BY A33- RELATED ANTIGENS
; FILE REFERENCE: P1216R1 (US)
; CURRENT APPLICATION NUMBER: US/09/953,499
; CURRENT FILING DATE: 2001-09-14
; PRIOR APPLICATION NUMBER: US/09/254,465
; PRIOR FILING DATE: 1999-03-05
; PRIOR APPLICATION NUMBER: PCT/US98/24855
; PRIOR FILING DATE: 1998-11-20
; PRIOR APPLICATION NUMBER: US 60/066,364
; PRIOR FILING DATE: 1997-11-21
; PRIOR APPLICATION NUMBER: US 60/078,936
; PRIOR FILING DATE: 1998-03-20
; PRIOR APPLICATION NUMBER: PCT/US98/19437
; PRIOR FILING DATE: 1998-09-17
; NUMBER OF SEQ ID NOS: 30
; SEQ ID NO 2
; LENGTH: 321
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-953-499-2

Query Match 100.0%; Score 1688; DB 4; Length 321;
Best Local Similarity 100.0%; Pred. No. 1.4e-165;
Matches 321; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MGILLGLLLGLHGLTVDYGRPILEVPESVTGPKGKDVNLPCYDPLQGYTQVLVKWLVR 60
Db 1 MGILLGLLLGLHGLTVDYGRPILEVPESVTGPKGKDVNLPCYDPLQGYTQVLVKWLVR 60

QY 61 GSDPVTIFLRDSSGDHIQQAQYQGRHVS HKVPGDVSLQSLTLEMDRSHYTCVETWQTP 120
Db 61 GSDPVTIFLRDSSGDHIQQAQYQGRHVS HKVPGDVSLQSLTLEMDRSHYTCVETWQTP 120

QY 121 DGNQVVRDKITELRVOKLSVSKPTVTGSGYGTVPQGMRIISLQCCARGSPPISYIWKQ 180
Db 121 DGNQVVRDKITELRVOKLSVSKPTVTGSGYGTVPQGMRIISLQCCARGSPPISYIWKQ 180

QY 181 QTNNOEPIKVATLSTLLFPKPAVIADSGSYFCTAKQGVSEQHSDIVKFFVKDSSKLLKTK 240
Db 181 QTNNOEPIKVATLSTLLFPKPAVIADSGSYFCTAKQGVSEQHSDIVKFFVKDSSKLLKTK 240

QY 241 TEAPTTMTYPLKATSTVKQSDWTTMDGVLGETSAGPGKSLPVFAIILISLCCMWVFT 300
Db 241 TEAPTTMTYPLKATSTVKQSDWTTMDGVLGETSAGPGKSLPVFAIILISLCCMWVFT 300

QY 301 MAYIMLCRKTSQOEHVYEAAAR 321
Db 301 MAYIMLCRKTSQOEHVYEAAAR 321

RESULT 3
US-09-369-247-63
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; Sequence 63, Application US/09369247
; Patent No. 656992
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: 44 Human Secreted Proteins
; FILE REFERENCE: P2024P1
; CURRENT APPLICATION NUMBER: US/09/369,247
; CURRENT FILING DATE: 1999-08-05
; EARLIER APPLICATION NUMBER: 60/074,118
; EARLIER FILING DATE: 1998-02-09
; EARLIER APPLICATION NUMBER: 60/074,157
; EARLIER FILING DATE: 1998-02-09
; EARLIER APPLICATION NUMBER: 60/074,137
; EARLIER FILING DATE: 1998-02-09
; EARLIER APPLICATION NUMBER: 60/074,341
; EARLIER FILING DATE: 1998-02-09
; EARLIER APPLICATION NUMBER: 60/074,141
; EARLIER FILING DATE: 1998-02-09
; NUMBER OF SEQ ID NOS: 172
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 63
; LENGTH: 306
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (306)
; OTHER INFORMATION: Xaa equals stop translation
US-09-369-247-63

Query Match 67.4%; Score 1137; DB 4; Length 306;
Best Local Similarity 70.4%; Pred. No. 9.4e-109;
Matches 226; Conservative 0; Mismatches 1; Indels 94; Gaps 1;

QY 1 MGILLGLLLGLHGLTVDYGRPILEVPESVTGPKGKDVNLPCYDPLQGYTQVLVKWLVR 60
Db 1 MGILLGLLLGLHGLTVDYGRPILEVPESVTGPKGKDVNLPCYDPLQGYTQVLVKWLVR 60

QY 61 GSDPVTIFLRDSSGDHIQQAQYQGRHVS HKVPGDVSLQSLTLEMDRSHYTCVETWQTP 120
Db 61 GSDPVTIFLRDSSGDHIQQAQYQGRHVS HKVPGDVSLQSLTLEMDRSHYTCVETWQTP 120

QY 121 DGNQVVRDKITELRVOKLSVSKPTVTGSGYGTVPQGMRIISLQCCARGSPPISYIWKQ 180
Db 121 DGNQVVRDKITELRVQ----- 136

QY 181 QTNNOEPIKVATLSTLLFPKPAVIADSGSYFCTAKQGVSEQHSDIVKFFVKDSSKLLKTK 240
Db 137 -----KHSSKLLKTK 146

QY 241 TEAPTTMTYPLKATSTVKQSDWTTMDGVLGETSAGPGKSLPVFAIILISLCCMWVFT 300
Db 147 TEAPTTMTYPLKATSTVKQSDWTTMDGVLGETSAGPGKSLPVFAIILISLCCMWVFT 206

QY 301 MAYIMLCRKTSQOEHVYEAAAR 321
Db 207 MAYIMLCRKTSQOEHVYEAAAR 227

RESULT 4
US-09-763-902B-6
; Sequence 6, Application US/09763902B
; Patent No. 6838241
; GENERAL INFORMATION:
; APPLICANT: INCYTE PHARMACEUTICALS, INC.
; APPLICANT: TANG, Y. Tom
; APPLICANT: LAL, Preeti
; APPLICANT: BANDMAN, Olga
; APPLICANT: YUE, Henry
; APPLICANT: CORLEY, Neil C.
; APPLICANT: GUEGLER, Karl J.
; APPLICANT: GORGONE, Gina A.
; APPLICANT: BAUGHN, Mariah R.
```

```
; APPLICANT: PATTERSON, Chandra
; TITLE OF INVENTION: PROTEIN TRANSPORT-ASSOCIATED MOLECULES
; FILE REFERENCE: PF-0577 PCT
; CURRENT APPLICATION NUMBER: US/09/763,902B
; CURRENT FILING DATE: 2002-08-22
; PRIOR APPLICATION NUMBER: 60/098,206
; PRIOR FILING DATE: 1998-08-27
; NUMBER OF SEQ ID NOS: 16
; SOFTWARE: PERL Program
; SEQ ID NO 6
; LENGTH: 175
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc feature
; OTHER INFORMATION: Incyte ID No. 68382411871275CD1
US-09-763-902B-6

Query Match      51.5%; Score 870; DB 4; Length 175;
Best Local Similarity 64.7%; Pred. No. 1.4e-81;
Matches 174; Conservative 0; Mismatches 1; Indels 94; Gaps 1;

QY 1 MGILGLLLGLHGLTVDTYGRPILEVPESVTGPKGDNVLPCTYDPLQGYTVLVKLVQR 60
DB 1 MGILGLLLGLHGLTVDTYGRPILEVPESVTGPKGDNVLPCTYDPLQGYTVLVKLVQR 60

QY 61 GSDPVTIFLRDSSGDHIQQAQYQGRHLVSHK-----VPGDVLSQLSTLEMDRSHYTCV 120
DB 61 GSDPVTIFLRDSSGDHIQQAQYQGRHLVSHKPGDVLSQLSTLEMDRSHYTCV 120

QY 121 DGNQVRDKITELRVQKLSVSKPTVTGSGYGTVPQGMRLSLOQCAR-GSPPIIS 180
DB 121 DGNQVRDKITELRVQ-----136

QY 181 QTNQPEIKVATLSTLLFKPAVIADSGSYFCTAKGVGSEQHSDIVKVVKDSKLLTK 240
DB 137 -----KHSSKLLTK 146

QY 241 TEAPTTMTYPLKATSVKQSDWTTMDMG 269
DB 147 TEAPTTMTYPLKATSVKQSDWTTMDMG 175

RESULT 5
US-09-188-930-189
; Sequence 189, Application US/09188930A
; Patent No. 6150502
; GENERAL INFORMATION:
; APPLICANT: Watson, James D.
; APPLICANT: Strachan, Lorna
; APPLICANT: Sleeman, Matthew
; APPLICANT: Onrust, Rene
; APPLICANT: Murison, James Greg
; TITLE OF INVENTION: Compositions Isolated From Skin Cells
; TITLE OF INVENTION: and Methods For Their Use
; FILE REFERENCE: 11000.1011c1
; CURRENT APPLICATION NUMBER: US/09/188,930A
; CURRENT FILING DATE: 1998-11-09
; NUMBER OF SEQ ID NOS: 348
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 189
; LENGTH: 299
; TYPE: PRT
; ORGANISM: Human
; FEATURE:
; NAME/KEY: UNSURE
; LOCATION: (247)...(247)
; NAME/KEY: UNSURE
; LOCATION: (289)...(289)
US-09-188-930-189

Query Match      10.6%; Score 178.5; DB 3; Length 299;
Best Local Similarity 26.5%; Pred. No. 7.8e-10;
Matches 65; Conservative 36; Mismatches 93; Indels 51; Gaps 11;
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Matches 65; Conservative 36; Mismatches 93; Indels 51; Gaps 11;
QY 1 MGILGLLLGLHGLTVDTYGRPILEVPESVTGPKGDNVLPCTYDPLQGYTVLVKLVQR 60
DB 17 LAILLCSLALGSVTVHS--SEPEVRIPEN-----NPVKLSLAY---SGFSSPRVW--- 62
QY 61 GSDPVTIFLRDSSGDHIQQAQYQGRHLVSHK-----VPGDVLSQLSTLEMDRSHYTCV 115
DB 63 -----KFDQGDTRLVLCYNNKITASYEDRVTFELPTGITFKSVTRE--DTGTVTCMV 111
QY 116 TWQTPDGNQVVRDKITELRVQKLSVSKPTVTGSGYGTVPQGMRLSLOQCAR-GSPPIIS 174
DB 112 SEEGNSYGEVKVKLVIL-----VPPSKPTVNIPS-----SATIGNRAVITCSQDGSPPSE 163
QY 175 YIWK-----QOTN-----NQEPIKVATLSTLLFKPAVIADSGSYFCTAKGVGSEQH 222
DB 164 YTFKDGIVMPTNPKSTRAFNSNVLPNTTGGELVFDPLSASDTGEYSCEARNGYGTPT 223
QY 223 SDIVK 227
DB 224 SNAVR 228

RESULT 6
US-09-188-930-331
; Sequence 331, Application US/09188930A
; Patent No. 6150502
; GENERAL INFORMATION:
; APPLICANT: Watson, James D.
; APPLICANT: Strachan, Lorna
; APPLICANT: Sleeman, Matthew
; APPLICANT: Onrust, Rene
; APPLICANT: Murison, James Greg
; TITLE OF INVENTION: Compositions Isolated From Skin Cells
; TITLE OF INVENTION: and Methods For Their Use
; FILE REFERENCE: 11000.1011c1
; CURRENT APPLICATION NUMBER: US/09/188,930A
; CURRENT FILING DATE: 1998-11-09
; NUMBER OF SEQ ID NOS: 348
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 331
; LENGTH: 299
; TYPE: PRT
; ORGANISM: Human
; FEATURE:
US-09-188-930-331

Query Match      10.6%; Score 178.5; DB 3; Length 299;
Best Local Similarity 26.5%; Pred. No. 7.8e-10;
Matches 65; Conservative 36; Mismatches 93; Indels 51; Gaps 11;
QY 1 MGILGLLLGLHGLTVDTYGRPILEVPESVTGPKGDNVLPCTYDPLQGYTVLVKLVQR 60
DB 17 LAILLCSLALGSVTVHS--SEPEVRIPEN-----NPVKLSLAY---SGFSSPRVW--- 62
QY 61 GSDPVTIFLRDSSGDHIQQAQYQGRHLVSHK-----VPGDVLSQLSTLEMDRSHYTCV 115
DB 63 -----KFDQGDTRLVLCYNNKITASYEDRVTFELPTGITFKSVTRE--DTGTVTCMV 111
QY 116 TWQTPDGNQVVRDKITELRVQKLSVSKPTVTGSGYGTVPQGMRLSLOQCAR-GSPPIIS 174
DB 112 SEEGNSYGEVKVKLVIL-----VPPSKPTVNIPS-----SATIGNRAVITCSQDGSPPSE 163
QY 175 YIWK-----QOTN-----NQEPIKVATLSTLLFKPAVIADSGSYFCTAKGVGSEQH 222
DB 164 YTFKDGIVMPTNPKSTRAFNSNVLPNTTGGELVFDPLSASDTGEYSCEARNGYGTPT 223
QY 223 SDIVK 227
DB 224 SNAVR 228

RESULT 7
US-09-462-270-2
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; Sequence 2, Application US/09462270
; Patent No. 6358707
; GENERAL INFORMATION:
; APPLICANT: SmithKline Beecham Corporation
; TITLE OF INVENTION: Human P11 Antigen: A Cell Surface
; TITLE OF INVENTION: Receptor Involved in Platelet Aggregation
; FILE REFERENCE: GH-70150US
; CURRENT APPLICATION NUMBER: US/09/462,270
; CURRENT FILING DATE: 2000-01-05
; PRIOR APPLICATION NUMBER: 60/052,186
; PRIOR FILING DATE: 1997-07-10
; NUMBER OF SEQ ID NOS: 4
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 2
; LENGTH: 299
; TYPE: PRT
; ORGANISM: HOMO SAPIENS
US-09-462-270-2

Query Match 10.6%; Score 178.5; DB 3; Length 299;
Best Local Similarity 26.5%; Pred.No. 7.8e-10;
Matches 65; Conservative 36; Mismatches 93; Indels 51; Gaps 11;

QY 1 MGILLGLLLGHLTVDYGRPILEVPESVTGPKWGDVNLPCYDPLQYQTQVLVKLWVQR 60
Db 17 LAILLCSLALGSVTVHS-SEPEVRIPEN-----NPVKLSLAY---SGFSSPRVEW---- 62
QY 61 GSDPVTIFLRDSSDHTQQAQYQGRHLVSHK-----VPGDVSLQLSTLEMDDRSHYTCV 115
Db 63 -----KFDQGDVTLRLVCYNNKHTASVEDRVTFLEGTITFKSVTRE--DTGTYTCW 111
QY 116 TWOTPDGNQVVRDKITELRQKLSVSKPTVTGSGYGTVPQGMRIQLQCOAR-GSPPIIS 174
Db 112 SEEGSGSYGEVKVQLVL-----VPPSKPTVNIPS-----SATIGNRAVLTCSEQDGSPPSE 163
QY 175 YIYWK-----QQTN-----NOBPIKAVLSTLLFKPAVITADSGSVFCTAKGVGSEQH 222
Db 164 YTFKDGIVMPTNPKSTRAFSNSVYVLNPTTGELVFDPLSASDTGEYSCARNGYGTPT 223
QY 223 SDIVK 227
Db 224 SNAVR 228

RESULT 8
US-09-254-465A-1
; Sequence 1, Application US/09254465A
; Patent No. 6410708
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Ashkenazi, Avi J.
; APPLICANT: Fong, Sherman
; APPLICANT: Goddard, Audrey
; APPLICANT: Gurney, Austin L.
; APPLICANT: Napier, Mary A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Wood, William I.
; TITLE OF INVENTION: COMPOUNDS, COMPOSITIONS AND METHODS FOR THE TREATMENT
; OF DISEASES CHARACTERIZED BY A33- RELATED ANTIGENS
; FILE REFERENCE: P1216R1(US)
; CURRENT APPLICATION NUMBER: US/09/254,465A
; CURRENT FILING DATE: 1999-03-05
; PRIOR APPLICATION NUMBER: PCT/US98/24855
; PRIOR FILING DATE: 1998-11-20
; PRIOR APPLICATION NUMBER: US 60/066,364
; PRIOR FILING DATE: 1997-11-21
; PRIOR APPLICATION NUMBER: US 60/078,936
; PRIOR FILING DATE: 1998-03-20
; PRIOR APPLICATION NUMBER: PCT/US98/19437
; PRIOR FILING DATE: 1998-09-17
; NUMBER OF SEQ ID NOS: 30
; SEQ ID NO 1
; LENGTH: 299

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FILE REFERENCE: 011.00221
CURRENT APPLICATION NUMBER: US/09/397,243D
CURRENT FILING DATE: 1999-09-16
PRIOR APPLICATION NUMBER: 60/100,638
PRIOR FILING DATE: 1998-09-16
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patent in Ver. 2.1
SEQ ID NO 3
LENGTH: 299
TYPE: PRT
ORGANISM: Homo sapiens
US-09-397-243D-3

Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 7.8e-10;
Matches 65; Conservative 36; Mismatches 93; Indels 51; Gaps 11;

QY 1 MGIILGLLLHLTVDTYGRPILEVPEVSTGPKWGDVNLPCYDPLQGYTQVLVKLVQR 60
Db 17 LAILLCSLALGSGVTVHS-SEPEVRIPEN-----NPVKLSLAY---SGFSSPRVEW---- 62

QY 61 GSDPVTIFLRDSSGDHIQQAQYGRHLVSHK-----VPGDVSLLQSLTLEMDRSHVTCV 115
Db 63 -----KFDQDGTTLVLCVNNKITASYEDRVTFPLTGITFKSVTRE--DTGYTCMV 111

QY 116 TWQTPDGNQVVRDKITELRVQKLSVSKPTVTGSGYGVFTVPOGMRISLQCAR-GSPPTS 174
Db 112 SBEAGNSYGEVKVLIIVL-----VPPSKPTVNIIPS-----SATIGNRAVLTCSDQSGSPSE 163

QY 175 YIWK-----QQTN-----NOEPIKVAATLTLFKPAVIADSGSYFCTAGQGVSEQH 222
Db 164 YTFKDGIVMPTNPKSTRAFSNSVNLPTTGELVDFPLSDTGEYSCEARNGYGTPTMT 223

QY 223 SDIVK 227
Db 224 SNAVR 228

RESULT 15
US-09-906-700-119
Sequence 119 Application US/09906700
Patent No. 6723535
GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Ashkenazi, Avi
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerber, Hanspeter
APPLICANT: Gerritsen, Mary E.
APPLICANT: Goddard, A.
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kijavin, Ivar J.
APPLICANT: Mather, Jernie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas F.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/906,700
CURRENT FILING DATE: 2000-09-18

PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
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PRIOR APPLICATION NUMBER: PCT/US99/21547
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PRIOR APPLICATION NUMBER: PCT/US99/23089
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PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR FILING DATE: 1999-11-30
PRIOR APPLICATION NUMBER: PCT/US99/28564
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PRIOR APPLICATION NUMBER: PCT/US99/30911
PRIOR FILING DATE: 1999-12-20
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PRIOR APPLICATION NUMBER: PCT/US00/00219
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NUMBER OF SEQ ID NOS: 423
SEQ ID NO 119
LENGTH: 299
TYPE: PRT
ORGANISM: Homo sapiens
US-09-902-775A-119

Query Match 10.6%; Score 178.5; DB 4; Length 299;
Best Local Similarity 26.5%; Pred. No. 7.8e-10;
Matches 65; Conservative 36; Mismatches 93; Indels 51; Gaps 11;

QY 1 MGIILGLLLHLTVDTYGRPILEVPEVSTGPKWGDVNLPCYDPLQGYTQVLVKLVQR 60
Db 17 LAILLCSLALGSGVTVHS-SEPEVRIPEN-----NPVKLSLAY---SGFSSPRVEW---- 62

QY 61 GSDPVTIFLRDSSGDHIQQAQYGRHLVSHK-----VPGDVSLLQSLTLEMDRSHVTCV 115
Db 63 -----KFDQDGTTLVLCVNNKITASYEDRVTFPLTGITFKSVTRE--DTGYTCMV 111

QY 116 TWQTPDGNQVVRDKITELRVQKLSVSKPTVTGSGYGVFTVPOGMRISLQCAR-GSPPTS 174
Db 112 SBEAGNSYGEVKVLIIVL-----VPPSKPTVNIIPS-----SATIGNRAVLTCSDQSGSPSE 163

QY 175 YIWK-----QQTN-----NOEPIKVAATLTLFKPAVIADSGSYFCTAGQGVSEQH 222
Db 164 YTFKDGIVMPTNPKSTRAFSNSVNLPTTGELVDFPLSDTGEYSCEARNGYGTPTMT 223

QY 223 SDIVK 227
Db 224 SNAVR 228

RESULT 14
US-09-397-243D-3
Sequence 3, Application US/09397243D
Patent No. 6699688
GENERAL INFORMATION:
APPLICANT: Kornecki, Elizabeth
APPLICANT: Sobocka, Malgorzata B.
TITLE OF INVENTION: Human Platelet F11 Receptor

